

Supplementary Information

mRNA cap analogues substituted in the tetraphosphate chain with CX₂: identification of O-to-CCl₂ as the first bridging modification that confers resistance to decapping without impairing translation

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Supplementary figures and tables

Table 1 pK_a values of mononucleotide analogues **2a-2d** as determined by ³¹P NMR assay.

compound	concentration [mM]	pK _a ⁴ (gamma)*	pK _a ⁴ (beta)**	Mean pK _a ⁴
m ⁷ Gppp/NH ₄ ⁺ (2d)	15.0	6.55±0.02	6.62±0.04	6.56±0.03
m ⁷ GppCH ₂ p/Na ⁺ (2a)	5.49	8.63±0.02	8.63±0.01	8.63±0.01
m ⁷ GppCCl ₂ p/NH ₄ ⁺ (2b)	5.33	7.39±0.02	7.25±0.02	7.32±0.07
m ⁷ GppCF ₂ p/NH ₄ ⁺ (2c)	7.50	6.21±0.02	6.29±0.03	6.25±0.04

* pK_a⁴ as determined using change of chemical shift for gamma-phosphate.

** pK_a⁴ as determined using change of chemical shift for beta-phosphate.

pK_a⁴ using change of chemical shift for alpha-phosphates was not determined due to only small changes in chemical shifts for alpha-phosphates upon pH change. The mean pK_a⁴ values were determined as weighted mean of pK_a⁴ (gamma) and pK_a⁴ (beta) values.

The collected data were fitted to DoseResp function in OriginLab software using following equation:

$$\text{chemical shift} = A_1 + \frac{(A_2 - A_1)}{1 + 10^{(pK_a - pH) * p}} \quad (1a)$$

where the **A**₁ and **A**₂ are bottom and top asymptote respectively, **p** is the hill slope. The pK_a values were determined for both gamma and beta phosphate and the weighted average with uncertainties were calculated using equations 2_{a-c}.

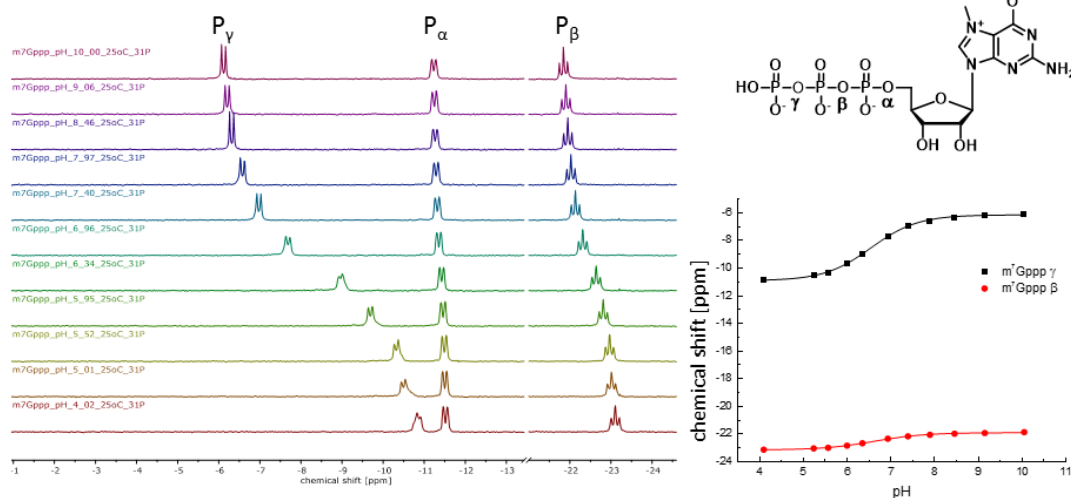
$$\bar{x}_w = \frac{\sum_{i=1}^N \frac{x_i}{u_i^2}}{\sum_{i=1}^N \frac{1}{u_i^2}} \quad (2a)$$

$$u_{int} = \sqrt{\sum_{i=1}^N \frac{1}{u_i^2}} \quad (2b)$$

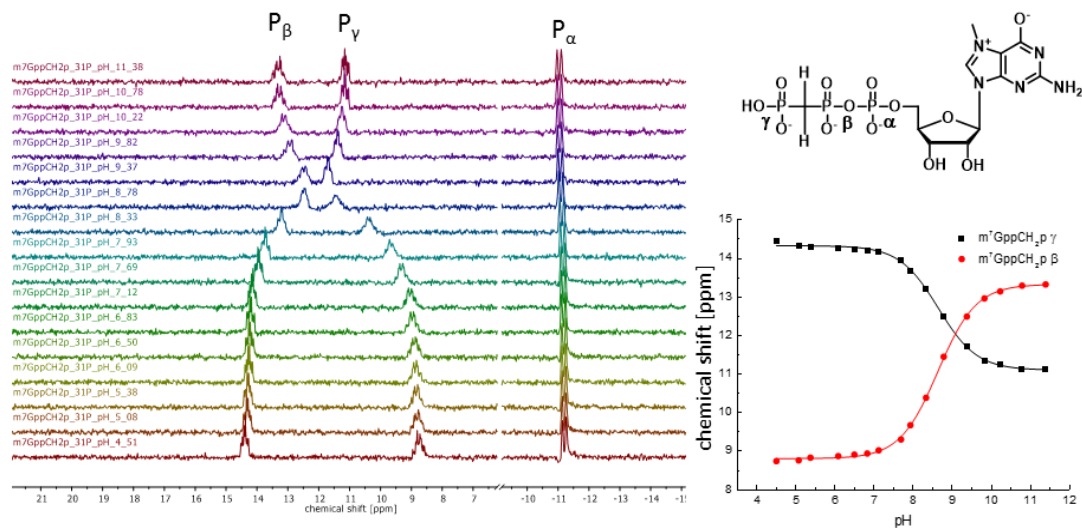
$$u_{ext} = \sqrt{\frac{u_{int}^2}{N-1} \sum_{i=1}^N \left(\frac{x_i - \bar{x}_w}{u_i} \right)^2} \quad (2c)$$

Fig. S1 Titrations of cap analogues **2a-2d** as monitored by ^{31}P NMR. The NMR spectra and titration curves for analogues **2d** (A), **2a** (B), **2b** (C) and **2c** (D).

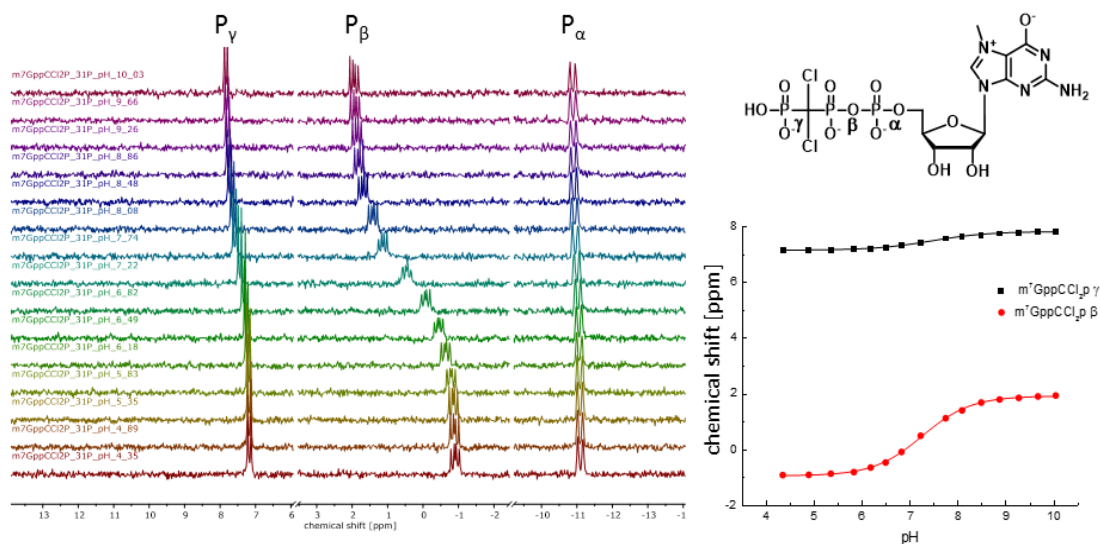
A



B



C



D

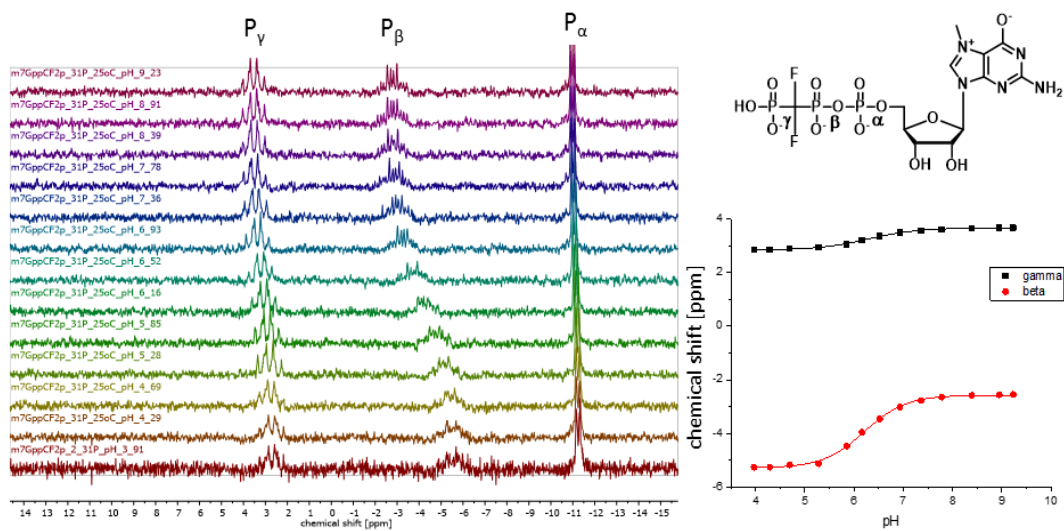


Table S2 Comparison for pK_a^4 values of mononucleoside triphosphates.

compound	pK_a^4 value*	compound	pK_a^4 value	compound	pK_a^4 value
Appp	7.1	Gppp	n.d.	m^7 Gppp	6.56 ± 0.03
AppCH ₂ p	8.4	GppCH ₂ p	n.d.	m^7 GppCH ₂ p	8.63 ± 0.01
AppCCl ₂ p	7.0	GppCCl ₂ p	7.49 ± 0.03	m^7 GppCCl ₂ p	7.32 ± 0.07
AppCF ₂ p	7.1	GppCF ₂ p	6.51 ± 0.07	m^7 GppCF ₂ p	6.25 ± 0.04

*Data from Blackburn et al.¹

Fig. S2 Titrations of GppCCl₂p (**4b**) and GppCF₂p (**4c**) as monitored by ³¹P NMR. The NMR spectra and titration curves for analogues **4b** (A) and **4c** (B).

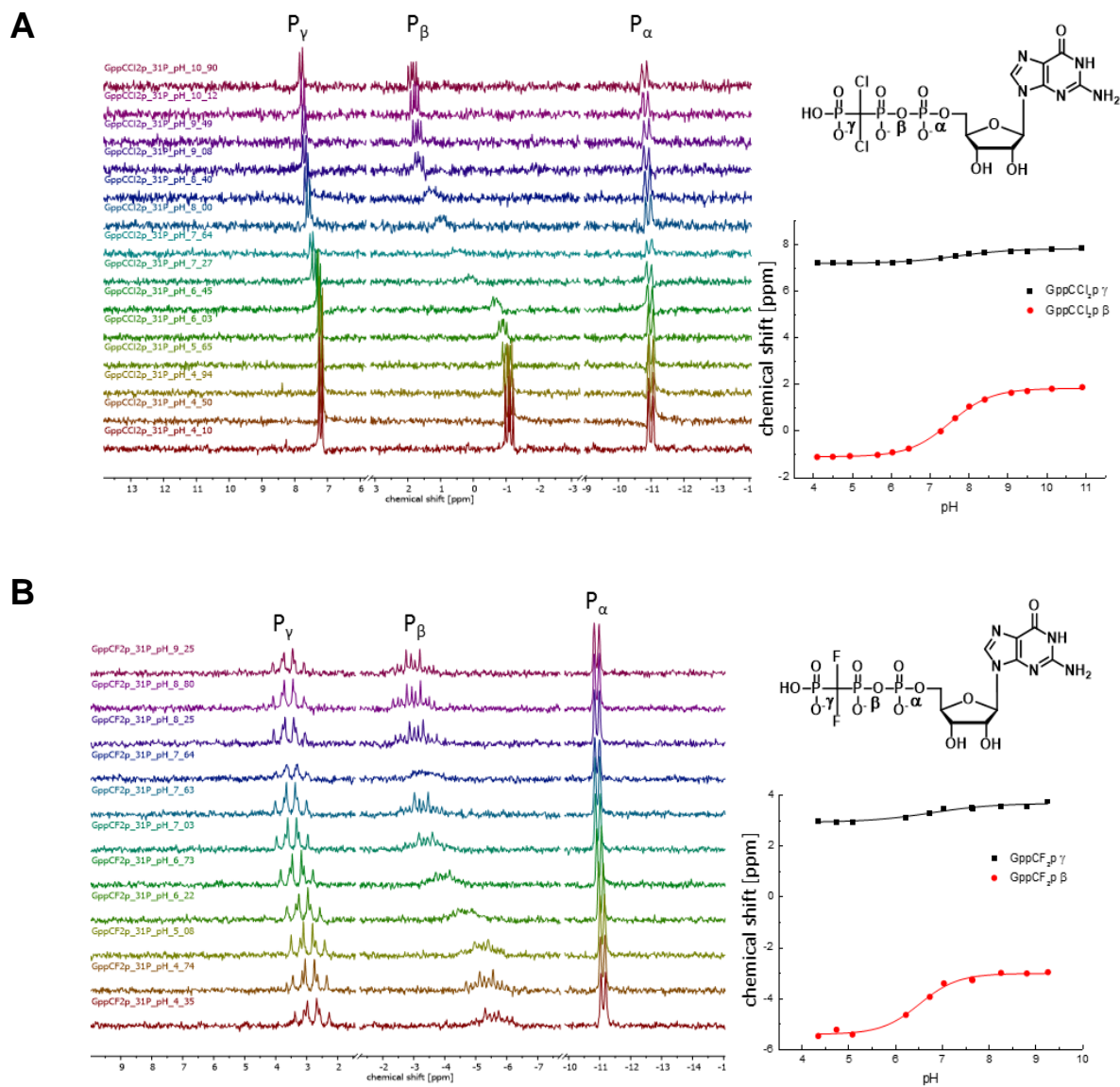


Fig. S3 Fluorescence titration curves for binding of cap analogues **2b-c** and **6b-c** to eIF4E.

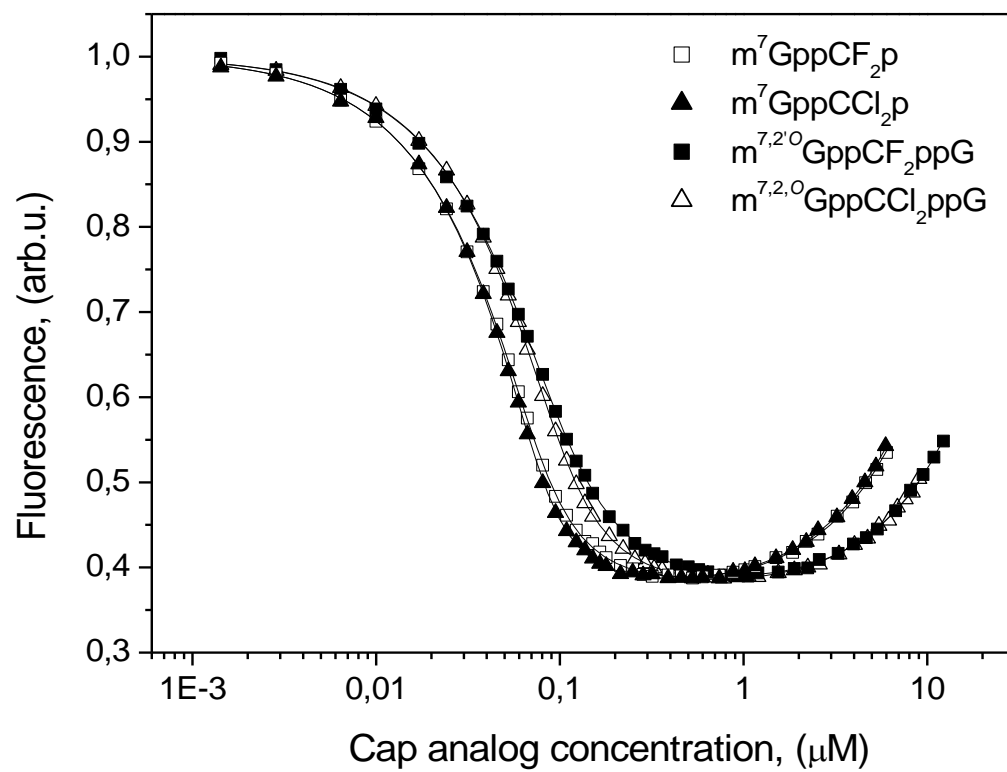


Fig. S4 DcpS and Dcp2 catalysed hydrolysis of various nucleotide substrates.

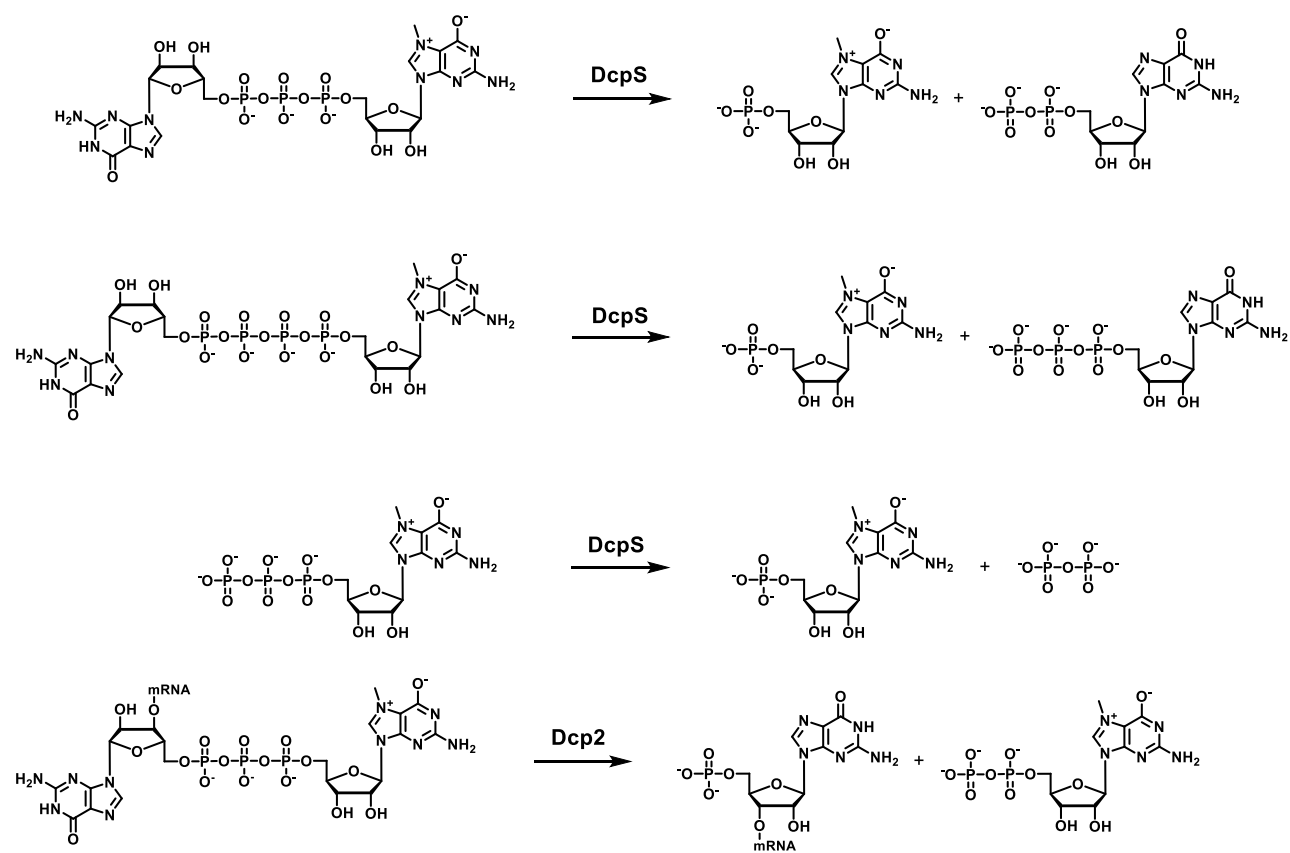


Table S3 Hydrolysis of cap analogues by human DcpS as monitored by HPLC. Data are average from duplicate experiments.

Cap analogue	% hydrolysis in given time			
	15 min	30 min	60 min	120 min
m ⁷ GpppG	67.0 ± 7.0	95.1 ± 3.0	98.2 ± 1.8	100 ± 0
m ⁷ Gppp (2d)	12.1 ± 0.9	22.3 ± 1.6	43.9 ± 3.5	77.6 ± 1.6
m ⁷ GppCH ₂ p (2a)	0	0	0	0
m ⁷ GppCCl ₂ p (2b)	0.2 ± 0.2	0.7 ± 0.0	1.1 ± 0.3	1.7 ± 0.7
m ⁷ GppCF ₂ p (2c)	2.9 ± 0.2	4.0 ± 0.8	6.9 ± 0.0	14.4 ± 0.6
m ⁷ GppppG (6d)	31.0 ± 0.8	55.7 ± 0.8	90.0 ± 2.3	100 ± 0
m ⁷ GppCH ₂ ppG (6a)	0	0	0	0
m ⁷ GppCCl ₂ ppG (6b)	0	0	0	0
m ⁷ GppCF ₂ ppG (6c)	13.7 ± 0.2	26.2 ± 0.4	50.1 ± 1.8	87.5 ± 0.4
m ₂ ^{7,2'-O} GppCCl ₂ ppG (7b)	0	0	0	0
m ₂ ^{7,2'-O} GppCF ₂ ppG (7c)	2.3 ± 0.4	4.7 ± 0.2	8.1 ± 0.7	15.7 ± 1.9

m₂^{7,2'-O}GppCH₂ppG (**7a**) was found to be resistant towards enzymatic hydrolysis by hDcpS in a similar assay conditions.² Error corresponds to standard deviation of two independent experimental points.

Fig. S5 Time course of DcpS mediated hydrolysis of modified cap analogues. Data are average from duplicate experiments.

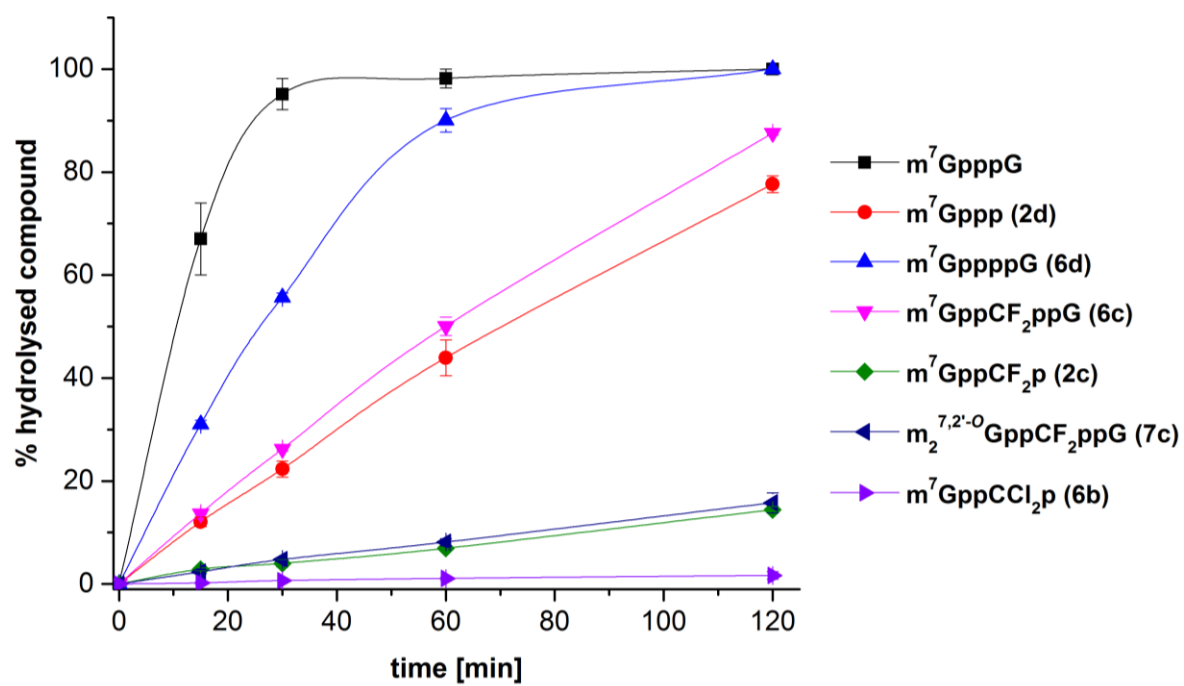
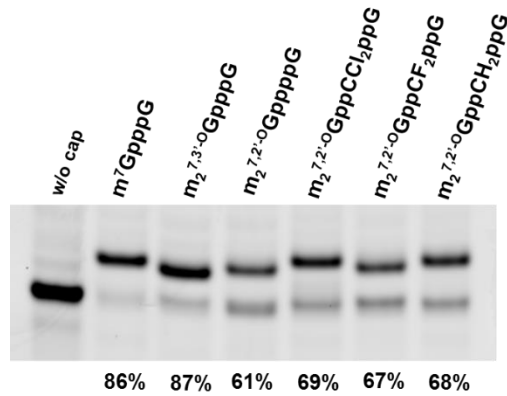


Fig. S6 Capping efficiencies (A) and susceptibilities (B) of 26-nt transcripts capped with various cap analogs to the hDcp2. Reactions were terminated at the indicated time points followed by denaturing PAGE, stained with SYBR Gold and visualized. Figure presents representative result of one biological repetition.

A



B

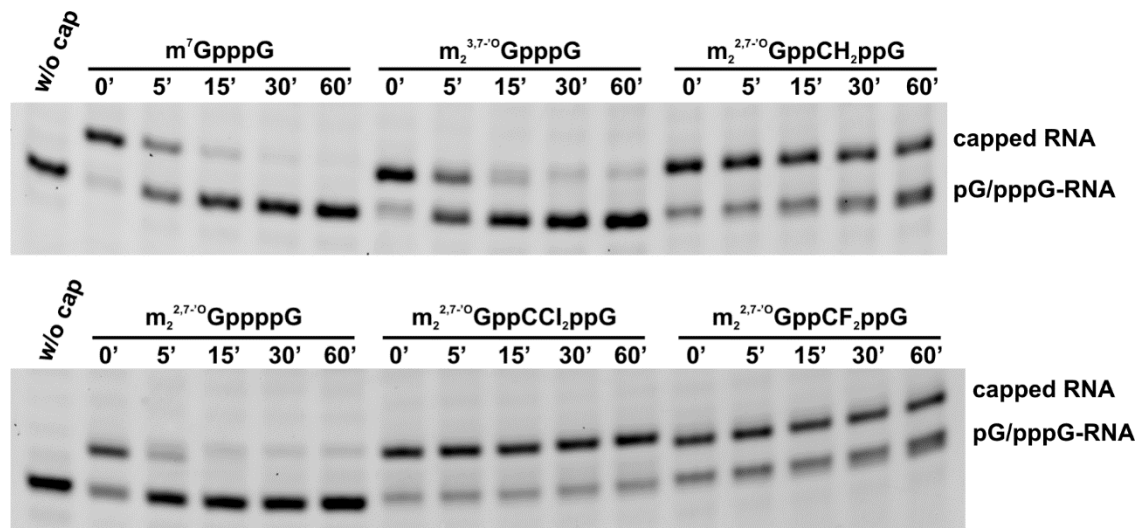
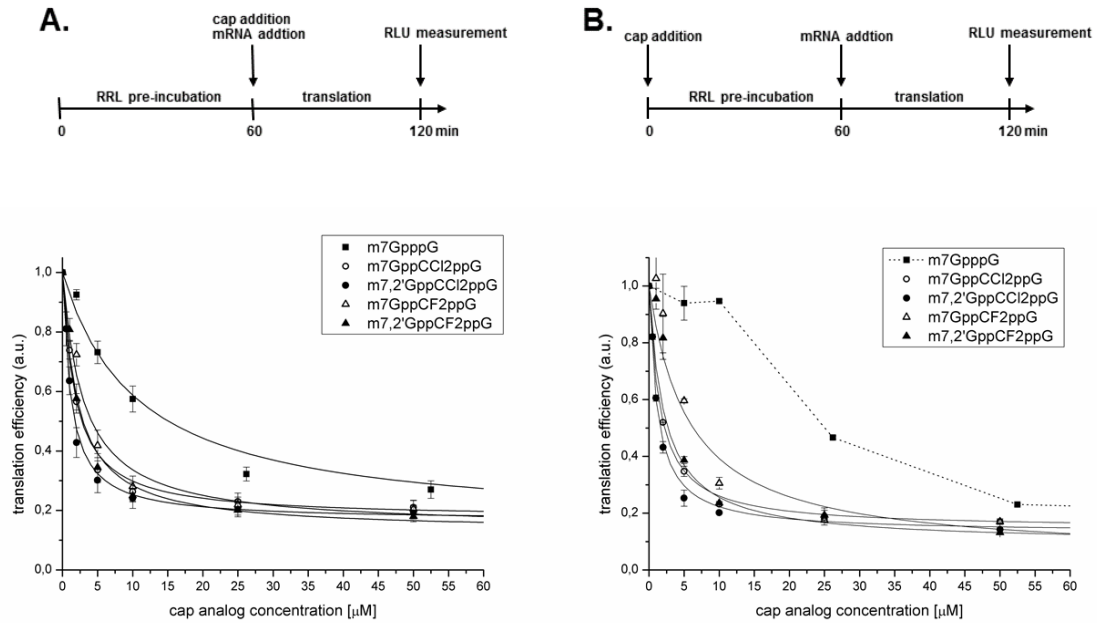
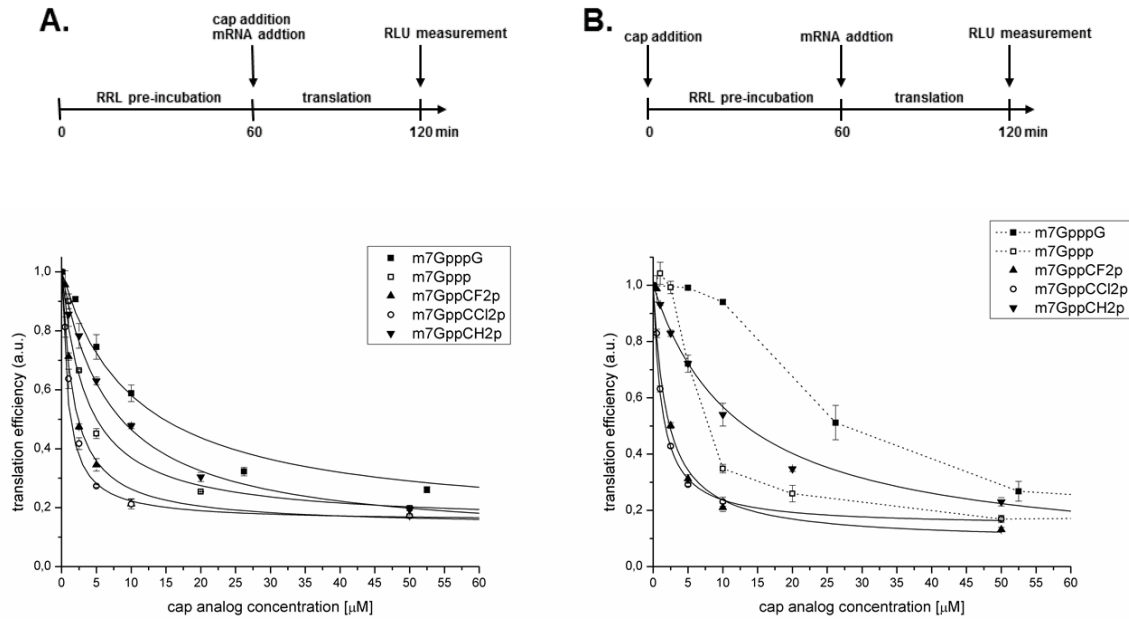


Fig. S7 Inhibition of translation of $m_2^{7,3'-O}$ GpppG-capped luciferase encoding mRNA in rabbit reticulocyte lysate (RRL) by dinucleotide cap analogues.



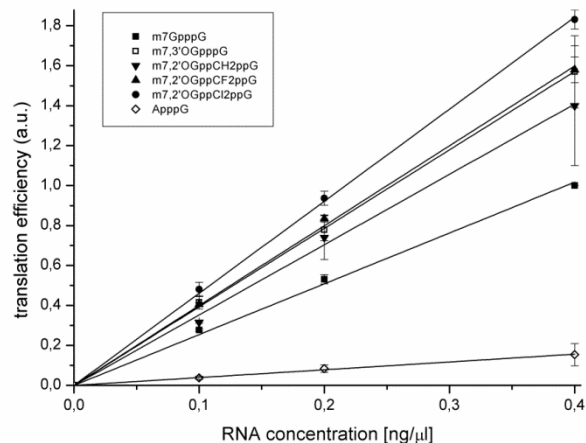
Inhibition of translation of $m_2^{7,3'-O}$ GpppG-capped luciferase encoding mRNA in rabbit reticulocyte lysate (RRL) by m^7 GpppG, m^7 GppCF₂pG, $m_2^{7,2'-O}$ GppCF₂pG, m^7 GppCl₂pG and $m_2^{7,2'-O}$ GppCl₂pG. In experiment (A) the cap analogue and luciferase mRNA were added to RRL at the same time point. In experiment (B), to test stability of presented here cap analogues in reticulocyte lysate, the cap analogue was preincubated for 1 hour in RRL prior to addition of mRNA and start of translation. As it is seen in graph (B), the inhibitory properties of unmodified m^7 GpppG is significantly diminished upon incubation in RRL (dotted line). In both experiments the luciferase activity was measured after 1 hour after mRNA addition. In the figure are shown data of 3 independent inhibition experiments (\pm SE) and 2 stability experiments (\pm SE), normalized to the translation efficiency without cap analog.

Fig. S8 Inhibition of translation of $m_2^{7,3'-O}$ GpppG-capped luciferase encoding mRNA in rabbit reticulocyte lysate (RRL) by mononucleotide cap analogues.



Inhibition of translation of $m_2^{7,3'-O}$ GpppG-capped luciferase encoding mRNA in rabbit reticulocyte lysate (RRL) by m^7 GpppG, m^7 GTP, m^7 GppCF₂p, m^7 GppCl₂p and m^7 GppCH₂p. In experiment (A) the cap analogue and luciferase mRNA were added to RRL at the same time point. In experiment (B), to test stability of presented here cap analogues in reticulocyte lysate, the cap analogue was preincubated for 1 hour in RRL prior to addition of mRNA and start of translation. As it is seen in graph (B), the inhibitory properties of unmodified m^7 GpppG and m^7 GTP are significantly diminished upon incubation in RRL (dotted line). In both experiments the luciferase activity was measured after 1 hour after mRNA addition. In the figure are shown data of 3 independent inhibition experiments (\pm SE) and 2 stability experiments (\pm SE), normalized to the translation efficiency without cap analog.

Fig. S9 Translation efficiencies of mRNA encoding firefly luciferase capped with cap analogues **7a-7c**.



Translation of mRNAs encoding firefly luciferase capped with $m_2^{7,2'-O}$ GppCF₂pG, $m_2^{7,2'-O}$ GppCl₂pG and $m_2^{7,2'-O}$ GppCH₂pG, and with standard caps: m⁷GpppG and $m_2^{7,3'-O}$ GpppG (Anti Reverse Cap Analog). Activity of luciferase (in RLU) synthesized in rabbit reticulocyte lysate were normalized to the activity obtained with m⁷GpppG-capped RNA at the highest concentration used. The mean of 3 independent translation experiments (\pm SE) and showed in the graph as a function of capped luciferase mRNA concentration (except $m_2^{7,2'-O}$ GppCF₂pG with 2 experiments). A transcript capped with non-functional ApppG dinucleotide was added as a control of cap-dependent translation in RRL. After linear fitting to the experimental data points the slope values (of the linear regression equation) calculated for the differentially capped luciferase RNAs were compared to the slope value obtained for m⁷GpppG-capped luciferase RNA that was set as = 1.

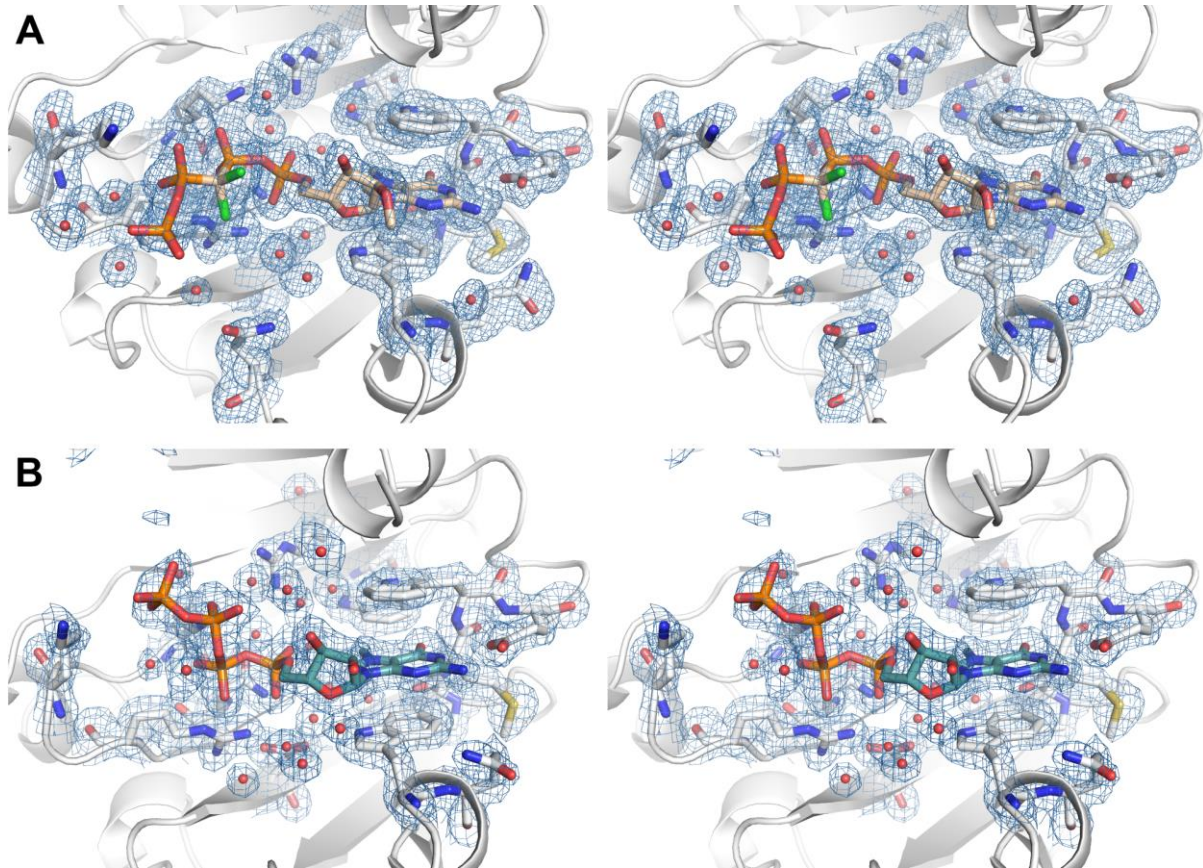
Table S4. Data collection and refinement statistics.

	eIF4E / 6d	eIF4E / 7b
PDB ID	5J5O	5J5Y
Wavelength	0.9184	0.9184
Resolution range	36.82–1.867 (1.934–1.867)	37–1.748 (1.81–1.748)
Space group	P 1	P 1
Unit cell	37.99 38.03 146.64 88.553 84.704 76.667	38.04 38.06 146.74 88.356 95.626 103.544
Total reflections	156370 (10888)	194484 (14491)
Unique reflections	60259 (5024)	72988 (6391)
Multiplicity	2.6 (2.2)	2.7 (2.3)
Completeness (%)	0.91 (0.76)	0.91 (0.78)
Mean I/sigma(I)	10.46 (2.04)	12.26 (2.46)
Wilson B-factor	19.87	20.00
R-merge	0.07081 (0.5346)	0.05243 (0.3991)
R-meas	0.08663 (0.6678)	0.06398 (0.499)
CC1/2	0.995 (0.677)	0.997 (0.825)
CC*	0.999 (0.898)	0.999 (0.951)
Reflections used in refinement	60257 (5024)	72986 (6391)
Reflections used for R-free	2234 (187)	3650 (320)
R-work	0.2209 (0.2829)	0.2364 (0.2772)
R-free	0.2637 (0.3532)	0.2860 (0.3429)
CC(work)	0.915 (0.832)	0.904 (0.824)
CC(free)	0.894 (0.733)	0.856 (0.700)
Number of non-hydrogen atoms	6401	5784
macromolecules	5880	5431
ligands	160	82

Protein residues	720	680
RMS(bonds)	0.008	0.007
RMS(angles)	1.03	0.95
Ramachandran favored (%)	96	95
Ramachandran allowed (%)	3.7	5.4
Ramachandran outliers (%)	0	0
Rotamer outliers (%)	1.9	1.6
Clashscore	6.70	5.77
Average B-factor	35.25	38.41
macromolecules	35.35	38.69
ligands	44.87	35.93
solvent	29.32	33.48
Number of TLS groups	33	33

Statistics for the highest-resolution shell are shown in parentheses.

Fig. S10 Wall-eye stereo view of eIF4E/7d cap-binding pocket with 2Fo-Fc electron density map contoured at 1.0 sigma.



A - Wall-eye stereo view of eIF4E/7b (PDB id: 5J5Y) cap-binding pocket with 2Fo-Fc electron density map contoured at 1.0 sigma; B - Wall-eye stereo view of eIF4E/6d (PDB id: 5J5O) cap-binding pocket with 2Fo-Fc electron density map contoured at 1.0 sigma.

Supplementary information – synthesis

General information

Reagents were purchased from Sigma-Aldrich and used without further purification, unless otherwise stated. Water used in the experiments was double distilled using MiliQ Milipore apparatus. Acetone was distilled over phosphorous pentoxide, triethylamine was distilled over potassium hydroxide and tetrahydrofuran was distilled over sodium. Dimethylformamide, dimethylsulfoxide and trimethylphosphate were kept over 4Å molecular sieves.

The nucleotides were purified by ion-exchange chromatography on a DEAE-Sephadex A-25 (HCO₃⁻ form) column. A column was loaded with the reaction mixture and washed thoroughly with water (until the eluate did not precipitate with AgNO₃ solution) to elute all material that does not bind to the resin. Then, the nucleotides were eluted using a linear gradient of triethylammonium hydrogen carbonate (TEAB) in deionized water. Fractions were analyzed spectrophotometrically at 260 nm and those containing the desired product were analyzed by reverse-phase HPLC and combined. After evaporation under reduced pressure with the repeated addition of ethanol to decompose TEAB, compounds were isolated as triethylammonium (TEA) salt. Yields were calculated on the basis of either sample weight or, preferably, optical density milliunits (mOD) of the product. Optical measurements for m7G mononucleotides were performed in 0.1 M phosphate buffer pH = 6 at 260 nm assuming $\epsilon_{260} = 11400 \text{ cm}^{-1} \text{ M}^{-1}$ for calculations. For guanine nucleotides and dinucleotide cap analogs measurements were conducted in 0.1 M phosphate buffer pH = 7 at 260 nm, assuming $\epsilon_{260} = 12080 \text{ cm}^{-1} \text{ M}^{-1}$ and $\epsilon_{260} = 22600 \text{ cm}^{-1} \text{ M}^{-1}$, respectively.

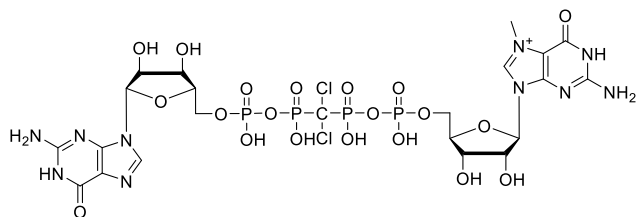
Analytical RP HPLC was performed with a Series 1200 instrument from Agilent Technologies on a Supelcosil LC-18-T HPLC column (4.6 x 250 mm, flow rate 1.3 mL min⁻¹) with a 0-25% linear gradient of methanol in 0.05 M ammonium acetate buffer (pH 5.9) for 15 min. Absorbance was monitored at 260 nm, while fluorescence was recorded at an excitation wavelength of 260 nm and an emission wavelength of 370 nm. Semi-preparative HPLC was performed on the same apparatus equipped with a Discovery Reverse-Phase Amide C-16 HPLC column (25 cm x 21.2 mm, 5 µm, flow rate 5.0 mL min⁻¹) and UV detection at 254 nm. The purity and homogeneity of each final product were confirmed by RP HPLC, high resolution mass spectrometry HRMS (ES⁻) and ¹H NMR and ³¹P NMR spectroscopy. Mass spectra were recorded with a high resolution LTQ Orbitrap Velos (Thermo Scientific). NMR spectra were recorded at 25 °C with a Varian UNITY-plus spectrometer at 399.94 MHz (¹H NMR) and 161.90 MHz (³¹P NMR). All chemical shifts (δ) are given in ppm and coupling constants (J) are given in Hz. ¹H NMR chemical shifts were calibrated to sodium 3-trimethylsilyl-[2,2,3,3-D₄]-propionate (TSP) in D₂O as an internal standard. ³¹P NMR chemical shifts were reported to 20% phosphorus acid in D₂O as an external standard. The raw NMR files were processed using ACD/Labs 12.0 Software.

Synthesis of previously described compounds

Methylenebisphosphonate containing analogues: m^7GppCH_2ppG (**6a**)¹ and $m_2^{7,2'-O}GppCH_2ppG$ (**7a**)¹, imidazolide derivatives: $m^7Gmp-Im$ (**1**)³, $Gmp-Im$ (**3**)⁴ and $m_2^{7,2'-O}Gmp-Im$ (**5**)⁴ and analogues unmodified in the polyphosphate bridge: $m^7GppppG$ (**6d**)⁵, $m_2^{7,2'-O}GppppG$ (**7d**)⁴, $m_2^{7,3'-O}GppppG$, m^7GpppG ⁵ and m^7Gppp (**2d**) were obtained as previously described.

Synthesis of m⁷GppCCl₂ppG

P1-(7-methylguanosin-5'-yl) P4-guanosin-5'-yl 2,3-dichloromethylenetetraphosphate

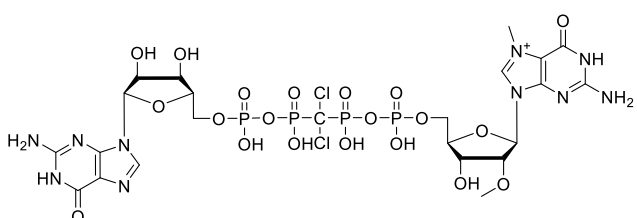


m⁷Gmp-Im (108 mg, 0.24 mmol) and GppCCl₂p (TEA salt, 85 mg, 0.095 mmol) were mixed in anhydrous DMF (4 ml) followed by addition of anhydrous ZnCl₂ (103 mg, 0.76 mmol). After 4 h reaction was

completed and quenched by addition of solution of EDTA (283 mg, 0.76 mmol) and NaHCO₃ (142 mg, 1.69 mmol) in water. Product was purified employing ion-exchange chromatography (Sephadex resin) and obtained as glassy solid (TEA salt, 1188 opt. u., 0.057 mmol, 60%). Further purification was performed using preparative HPLC yielding in 35.5 mg of final compound (NH₄⁺ salt, 0.035 mmol, 37%). **¹H NMR** (400 MHz, D₂O): δ 9.20 (1 H, s, H8_{m7G}), 8.05 (1 H, s, H8_G), 5.98 (1H, d, *J*_{1'-2'} = 3.3 Hz, H1'_{m7G}), 5.84 (1H, d, *J*_{1'-2'} = 6.2 Hz, H1'_G), 4.73 (1H, dd, *J*_{1'-2'} = 6.2 Hz, *J*_{2'-3'} = 5.4 Hz, H2'_G), 4.64 (1H, dd, *J*_{1'-2'} = 3.3 Hz, *J*_{2'-3'} = 5.1 Hz, H2'_{m7G}), 4.54 (1H, m, H3'_{m7G}), 4.50 (1H, dd, *J*_{2'-3'} = 5.4 Hz, *J*_{3'-4'} = 4.4 Hz, H3'_G), 4.41-4.25 (6H, m; H4'_{m7G}, H4'_G, H5'_{m7G}, H5'_G, H5''_{m7G}, H5''_G), 4.08 (3H, s, CH₃); **³¹P NMR** (162 MHz, D₂O) δ -10.81 (2P; P_{α,δ}), -1.41 (2P; P_{β,γ}). **HRMS** (ESI⁻) calc. for C₂₂H₂₉Cl₂N₁₀O₂₀P₄⁻ requires 946.9893, found 946.9901.

Synthesis of m₂^{7,2'-O}GppCCl₂ppG

P1-(7, 2'-O-dimethylguanosin-5'-yl) P4-guanosin-5'-yl 2,3-dichloromethylenetetraphosphate

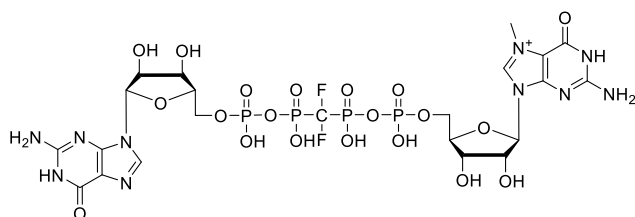


m₂^{7,2'-O}Gmp-Im (90 mg, 0.19 mmol) and GppCCl₂p (TEA salt, 57 mg, 0.064 mmol) were mixed in anhydrous DMF (3 ml) followed by addition of anhydrous ZnCl₂ (70 mg, 0.51 mmol). After 4 h reaction was completed and

quenched by addition of solution of EDTA (190 mg, 0.51 mmol) and NaHCO₃ (95 mg, 1.13 mmol) in water. Product was purified employing ion-exchange chromatography (Sephadex resin) and obtained as glassy solid (TEA salt, 900 opt. u., 0.043 mmol, 67%). Final compound was changed into sodium salt on Dowex (Na⁺ form) yielding 36.7 mg of final compound (0.035 mmol, 55%). **¹H NMR** (400 MHz, D₂O): δ 9.20 (1 H, s, H8_{m7G}), 8.76 (1 H, s, H8_G), 6.07 (1H, d, *J*_{1'-2'} = 3.1 Hz, H1'_{m7G}), 5.96 (1H, d, *J*_{1'-2'} = 4.6 Hz, H1'_G), 4.68 (1H, t, *J*_{1'-2'/2'-3'} = 4.8 Hz, H2'_G), 4.60 (1H, t, *J*_{2'-3'/3'-4'} = 5.3 Hz, H3'_{m7G}), 4.54 (1H, t, *J*_{2'-3'/3'-4'} = 4.4 Hz, H3'_G), 4.40-4.24 (6H, m; H4'_{m7G}, H4'_G, H5'_{m7G}, H5'_G, H5''_{m7G}, H5''_G), 4.10 (3H, s, CH₃), 3.58 (3H, s, OCH₃); **³¹P NMR** (162 MHz, D₂O) δ -10.78 (2P; P_{α,δ}), -1.35 (2P; P_{β,γ}). **HRMS** (ESI⁻) calc. for C₂₃H₃₁Cl₂N₁₀O₂₀P₄⁻ requires 961.0049, found 961.0056.

Synthesis of m⁷GppCF₂ppG

P1-(7-methylguanosin-5'-yl) P4-guanosin-5'-yl 2,3-difluoromethylenetetraphosphate

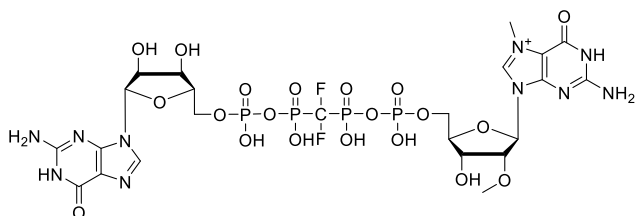


m⁷Gmp-Im (50 mg, 0.132 mmol) and GppCF₂p (ammonium salt, 55 mg, 0.088 mmol) were mixed in anhydrous DMF (2 ml) followed by addition of anhydrous ZnCl₂ (96 mg, 0.70 mmol). After 4 h reaction was

completed and quenched by addition of solution of EDTA (260 mg, 0.70 mmol) and NaHCO₃ (130 mg, 1.56 mmol) in water. Product was purified employing ion-exchange chromatography (Sephadex resin) and obtained as glassy solid (TEA salt, 920 opt. u., 0.044 mmol, 50%). Further purification was performed using preparative HPLC yielding in 17.7 mg of final compound (NH₄⁺ salt, 0.018 mmol, 20%). Final compound was changed into sodium salt on Dowex (Na⁺ form) yielding 17.9 mg of final compound (0.018 mmol, 20%). **¹H NMR** (400 MHz, D₂O): δ 9.12 (1 H, s, H8_{m7G}), 8.04 (1 H, s, H8_G), 5.98 (1H, d, *J*_{1'-2'} = 3.7 Hz, H1'_{m7G}), 5.85 (1H, d, *J*_{1'-2'} = 6.5 Hz, H1'_G), 4.74 (1H, t, *J*_{1'-2'/2'-3'} = 5.7 Hz, H2'_G), 4.65 (1H, t, *J*_{1'-2'/2'-3'} = 4.4 Hz, H2'_{m7G}), 4.52 (1H, dd, *J*_{2'-3'} = 5.0 Hz, *J*_{4'-3'} = 3.0 Hz, H3'_{m7G}), 4.49 (1H, t, *J*_{2'-3'/4'-3'} = 4.9 Hz, H3'_G), 4.19-4.43 (6H, m; H4'_{m7G}, H4'_G, H5'_{m7G}, H5'_G, H5''_{m7G}, H5''_G), 4.08 (3H, s, CH₃); **³¹P NMR** (162 MHz, D₂O) δ -11.08 (2P; P_{α,δ}), -6.37 (2P; P_{β,γ}). **HRMS** (ESI) calc. for C₂₂H₂₉F₂N₁₀O₂₀P₄⁻ requires 915.0484, found 915.0501.

Synthesis of m₂^{7,2'-O}GppCF₂ppG

P1-(7,2'-O-dimethylguanosin-5'-yl) P4-guanosin-5'-yl 2,3-difluoromethylenetetraphosphate

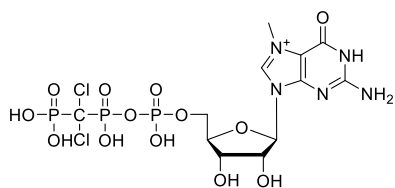


m^{7,2'-O}Gmp-Im (45 mg, 0.096 mmol) and GppCF₂p (ammonium salt, 40 mg, 0.064 mmol) were mixed in anhydrous DMF (1 ml) followed by addition of anhydrous ZnCl₂ (32 mg, 0.24 mmol). After 4 h reaction was

completed and quenched by addition of solution of EDTA (90 mg, 0.24 mmol) and NaHCO₃ (42 mg, 0.50 mmol) in water. Product was purified employing ion-exchange chromatography (Sephadex resin) and obtained as glassy solid (TEA salt, 1010 opt. u., 0.048 mmol, 55%). Further purification was performed using preparative HPLC yielding in 20.0 mg of final compound (NH₄⁺ salt, 0.020 mmol, 37%). Final compound was changed into sodium salt on Dowex (Na⁺ form) yielding 20.5 mg of final compound (0.021 mmol, 33%). **¹H NMR** (400 MHz, D₂O): δ 8.02 (1 H, s, H8_G), 6.01 (1H, br.s., H1'_{m7G}), 5.82 (1H, d, *J*_{1'-2'} = 6.7 Hz, H1'_G), 4.73 (1H, t, *J*_{1'-2'/2'-3'} = 5.6 Hz, H2'_G), 4.56 (1H, t, *J*_{2'-3'/3'-4'} = 5.4 Hz, H3'_{m7G}), 4.49-4.53 (1H, m, H3'_G), 4.20-4.43 (6H, m; H4'_{m7G}, H4'_G, H5'_{m7G}, H5'_G, H5''_{m7G}, H5''_G), 4.09 (3H, s, CH₃), 3.59 (3H, s, OCH₃); **³¹P NMR** (162 MHz, D₂O) δ -11.11 (2P; P_{α,δ}), -6.35 (2P; P_{β,γ}). **HRMS** (ESI) calc. for C₂₃H₃₁F₂N₁₀O₂₀P₄⁻ requires 929.0640, found 929.0656.

Synthesis of m⁷GppCCl₂p

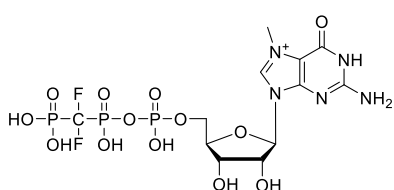
P1-(7-methylguanosin-5'-yl) 2,3-(dichloromethylene)triphosphate



To a suspension of dichlorobisphosphonate triethylammonium salt (400 mg, 1.20 mmol) in DMF (4 ml) anhydrous zinc chloride (164 mg, 1.21 mmol) was added, and the mixture was shaken until reagents dissolved. Then 7-methylguanosine imidazolidide (1700 opt. u., 0.15 mmol) was added, followed by addition of another portion of zinc chloride (164 mg, 1.21 mmol). The reaction was quenched after 4 hours by addition of water solution of EDTA (897 mg, 2.41 mmol) and NaHCO₃ (449 mg, 5.34 mmol). Product was purified employing ion-exchange chromatography on Sephadex resin in linear gradient from 0 to 1.1 M TEAB. Collected fractions were evaporated with several additions of ethanol yielding m⁷GppCCl₂p as triethylammonium salt (1020 opt.u., 0.089 mmol, 60%). Further purification was done by preparative HPLC. Collected fractions were freeze-dried several times until mass of the sample remained constant. Product was obtained as ammonium salt (26.6 mg, 0.041 mmol, 27%). **¹H NMR** (400 MHz, D₂O): δ 6.07 (1H, d, $J_{1'-2'}$ = 3.4 Hz, H1'), 4.70 (1H, dd, $J_{1'-2'}$ = 3.4 Hz, $J_{2'-3'}$ = 4.6 Hz, H2'), 4.58 (1H, dd, $J_{2'-3'}$ = 4.6 Hz, $J_{3'-4'}$ = 5.7 Hz, H3'), 4.43-4.30 (3H, m, H4', H5', H5''), 4.14 (3H, s, CH₃); **³¹P NMR** (162 MHz, D₂O) δ 7.90 (1P, d, J = 17.8 Hz, P γ), 0.88 (1P, dd, J = 17.5 Hz, J = 29.9 Hz, P β), -10.58 (1P, d, J = 30.3 Hz, P α). **HRMS** (ESI⁻) calc. for C₁₂H₁₇Cl₂N₅O₁₃P₃⁻ requires 601.9418, found 601.9425.

Synthesis of m⁷GppCF₂p

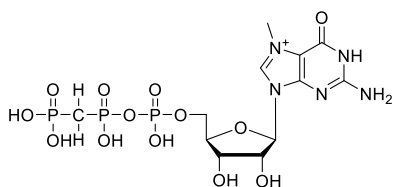
P1-(7-methylguanosin-5'-yl) 2,3-(difluoromethylene)triphosphate



To a suspension of difluorobisphosphonate triethylammonium salt (327 mg, 0.79 mmol) in DMF (4 ml) anhydrous zinc chloride (150 mg, 1.10 mmol) was added, and the mixture was shaken until reagents dissolved. Then 7-methylguanosine imidazolidide (1230 opt. u., 0.11 mmol) was added, followed by addition of another portion of zinc chloride (150 mg, 1.10 mmol). The reaction was quenched after 4 hours by addition of water solution of EDTA (821 mg, 2.21 mmol) and NaHCO₃ (410 mg, 5.34 mmol). Product was purified employing ion-exchange chromatography on Sephadex resin in linear gradient from 0 to 1.1 M TEAB. Collected fractions were evaporated with several additions of ethanol yielding m⁷GppCF₂p as triethylammonium salt (1032 opt.u., 0.091 mmol, 82%). Further purification was done by preparative HPLC. Collected fractions were freeze-dried several times until mass of the sample remained constant. Product was obtained as ammonium salt (31 mg, 0.049 mmol, 44%). **¹H NMR** (400 MHz, D₂O) δ ppm 9.22 (1H, s, H8), 6.08 (1H, d, $J_{1'-2'}$ = 3.7 Hz, H1'), 4.71 (1H, dd, $J_{1'-2'}$ = 3.7 Hz, $J_{2'-3'}$ = 4.7 Hz, H2'), 4.58 (1H, dd, $J_{2'-3'}$ = 4.7 Hz, $J_{3'-4'}$ = 5.5 Hz, H3'), 4.39 - 4.43 (1H, m, H4'), 4.35 (1H, m, H5'), 4.31 (1H, m, H5''), 4.14 (3H, s, CH₃). **³¹P NMR** (162 MHz, D₂O) δ 3.49 (1P, td, J = 75 Hz, J = 57 Hz, P γ), -3.51 (1P, m, P β), -10.90 (1P, d, J = 31.0 Hz, P α). **HRMS** (ESI⁻) calc. for C₁₂H₁₇F₂N₅O₁₃P₃⁻ requires 570.0009, found 570.0014.

Synthesis of m⁷GppCH₂p

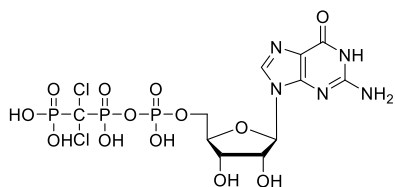
P1-(7-methylguanosin-5'-yl) 2,3-methylenetriphosphate



To a suspension of bisphosphonate triethylammonium salt (359 mg, 0.95 mmol) in DMF (4 ml) anhydrous zinc chloride (240 mg, 1.76 mmol) was added, and the mixture was shaken until reagents dissolved. Then 7-methylguanosine imidazolid (1248 opt. u., 0.11 mmol) was added, followed by addition of another portion of zinc chloride (240 mg, 1.76 mmol). The reaction was quenched after 4 hours by addition of water solution of EDTA (1.313 g, 3.53 mmol) and NaHCO₃ (657 mg, 7.82 mmol). Product was purified employing ion-exchange chromatography on Sephadex resin in linear gradient from 0 to 1.1 M TEAB. Collected fractions were evaporated with several additions of ethanol yielding m⁷GppCH₂p as triethylammonium salt (673 opt.u., 0.059 mmol, 54%). Product was changed into sodium salt on Dowex resin (Na⁺ form). **¹H NMR** (400 MHz, D₂O) δ ppm 5.96 (1H, d, $J_{1'-2'} = 3.6$ Hz, H1'), 4.56 (1H, dd, $J_{1'-2'} = 3.7$ Hz, $J_{2'-3'} = 4.9$ Hz, H2'), 4.42 (1H, t, $J = 5.2$ Hz, H3'), 4.29 (1H, dq, $J_{3'-4'} = 5.1$ Hz, $J_{4'-5'} = 2.6$ Hz, H4'), 4.23 (1H, m, H5'), 4.15 (1H, m, H5''), 4.01 (s, 3H), 2.21 (dd, $J = 20.8$, $J = 19.9$ Hz, 2H). **³¹P NMR** (162 MHz, D₂O) δ 15.03 (1P, d, $J = 8.8$ Hz, P γ), 10.21 (1P, dd, $J = 26.5$, $J = 8.9$, P β), -10.16 (1P, d, $J = 26.4$ Hz, P α). **HRMS** (ESI) calc. for C₁₂H₁₉N₅O₁₃P₃⁻ requires 534.0192, found 534.0195.

Synthesis of GppCCl₂p

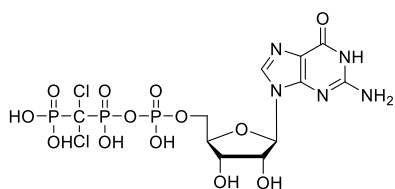
P1-guanosin-5'-yl-2,3-(dichloromethylene)triphosphate



To a suspension of dichlorobisphosphonate triethylammonium salt (720 mg, 1.61 mmol) in DMF (5 ml) anhydrous zinc chloride (218 mg, 1.61 mmol) was added, and the mixture was shaken until reagents dissolved. Then guanosine monophosphate imidazolid (200mg, 0.46 mmol) was added followed by addition of another portion of zinc chloride (218 mg, 1.61 mmol). The reaction was quenched after 4 hours by addition of water solution of EDTA (1120 mg, 3.2 mmol) and NaHCO₃ (590 mg, 7.04 mmol). Product was purified employing ion-exchange chromatography on Sephadex resin in linear gradient from 0 to 1.2 M TEAB. Collected fractions were evaporated with several additions of ethanol yielding GppCCl₂p as triethylammonium salt (370 mg, 0.41 mmol, 89%). **¹H NMR** (400 MHz, D₂O): δ 8.12 (1H, s, H8_G), 5.93 (1H, d, $J_{1'-2'} = 6.5$ Hz, H1'), 4.85 (1H, t, $J_{1'-2'/2'-3'} = 5.7$ Hz, H2'), 4.60 (1H, dd, $J_{2'-3'} = 5.2$ Hz, $J_{3'-4'} = 3.2$ Hz, H3'), 4.36 (1H, m, H4') 4.33-4.21 (2H, m, H5', H5''), 3.21 (q, $J = 7.5$ Hz, CH₂CH₃), 1.29 (t, $J = 7.3$ Hz, CH₂CH₃); **³¹P NMR** (162 MHz, D₂O) δ 7.71 (1P, d, $J = 17.8$ Hz, P γ), -0.65 (1P, dd, $J = 19.4$ Hz, $J = 30.1$ Hz, P β), -10.61 (1P, d, $J = 29.8$ Hz, P α). **HRMS** (ESI) calc. for C₁₁H₁₅Cl₂N₅O₁₃P₃⁻ requires 587.9262, found 587.9267.

Synthesis of GppCCl₂p

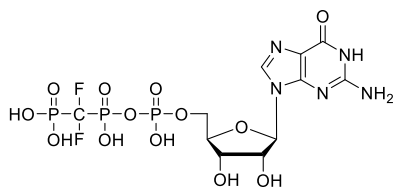
P1-guanosin-5'-yl-2,3-(dichloromethylene)triphosphate



To a suspension of dichlorobisphosphonate triethylammonium salt (720 mg, 1.61 mmol) in DMF (5 ml) anhydrous zinc chloride (218 mg, 1.61 mmol) was added, and the mixture was shaken until reagents dissolved. Then guanosine monophosphate imidazolidine (200mg, 0.46 mmol) was added followed by addition of another portion of zinc chloride (218 mg, 1.61 mmol). The reaction was quenched after 4 hours by addition of water solution of EDTA (1120 mg, 3.2 mmol) and NaHCO₃ (590 mg, 7.04 mmol). Product was purified employing ion-exchange chromatography on Sephadex resin in linear gradient from 0 to 1.2 M TEAB. Collected fractions were evaporated with several additions of ethanol yielding GppCCl₂p as triethylammonium salt (370 mg, 0.41 mmol, 89%). **¹H NMR** (400 MHz, D₂O): δ 8.12 (1H, s, H8_G), 5.93 (1H, d, *J*_{1'-2'} = 6.5 Hz, H1'), 4.85 (1H, t, *J*_{1'-2'/2'-3'} = 5.7 Hz, H2'), 4.60 (1H, dd, *J*_{2'-3'} = 5.2 Hz, *J*_{3'-4'} = 3.2 Hz, H3'), 4.36 (1H, m, H4') 4.33-4.21 (2H, m, H5', H5''), 3.21 (q, *J* = 7.5 Hz, CH₂CH₃), 1.29 (t, *J* = 7.3 Hz, CH₂CH₃); **³¹P NMR** (162 MHz, D₂O) δ 7.71 (1P, d, *J* = 17.8 Hz, Pγ), -0.65 (1P, dd, *J* = 19.4 Hz, *J* = 30.1 Hz, Pβ), -10.61 (1P, d, *J* = 29.8 Hz, Pα). **HRMS** (ESI) calc. for C₁₁H₁₅Cl₂N₅O₁₃P₃⁻ requires 587.9262, found 587.9267.

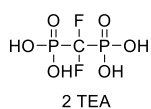
Synthesis of GppCF₂p

P1-guanosin-5'-yl-2,3-(difluoromethylene)triphosphate



To a suspension of dichlorobisphosphonate triethylammonium salt (700 mg, 1.70 mmol) in DMF (5 ml) anhydrous zinc chloride (218 mg, 1.61 mmol) was added, and the mixture was shaken until reagents dissolved. Then guanosine monophosphate imidazolidine (200mg, 0.46 mmol) was added followed by addition of another portion of zinc chloride (218 mg, 1.61 mmol). The reaction was quenched after 4 hours by addition of water solution of EDTA (1120 mg, 3.2 mmol) and NaHCO₃ (590 mg, 7.04 mmol). Product was purified employing ion-exchange chromatography on Sephadex resin in linear gradient from 0 to 1.2 M TEAB. Collected fractions were evaporated with several additions of ethanol yielding GppCF₂p as triethylammonium salt (224mg, 0.26 mmol, 57%). **¹H NMR** (400 MHz, D₂O): δ 8.30 (1H, s, H8_G), 5.86 (1H, d, *J*_{1'-2'} = 5.5 Hz, H1'), 4.62 (1H, t, *J*_{1'-2'/2'-3'} = 5.2 Hz, H2'), 4.45 (1H, dd, *J*_{2'-3'/3'-4'} = 4.2 Hz, H3'), 4.28 (1H, m, H4') 4.16 (2H, m, H5', H5''), 3.10 (q, *J* = 7.3 Hz, CH₂CH₃), 1.18 (t, *J* = 7.3 Hz, CH₂CH₃); **³¹P NMR** (162 MHz, D₂O) δ 7.71 (1P, td, *J* = 77.9×2, *J* = 58.8 Hz, Pγ), -4.08 (1P, m, Pβ), -10.63 (1P, d, *J* = 31.3 Hz, Pα). **HRMS** (ESI) calc. for C₁₁H₁₅F₂N₅O₁₃P₃⁻ requires 555.9853, found 555.98538.

Synthesis of methylenedifluorobisphosphonate

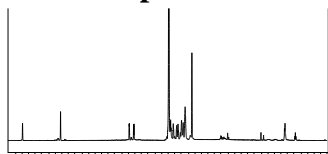


Tetraisopropyl methylenebisphosphonate (2.6 mL, 8.1 mmol) was placed in the oven dried two-neck roundbottom flask fitted with reflux condenser and flushed with argon. To the flask 20 mL of NaHMDS (1 M solution in THF, 20 mmol) was added and mixture was stirred for 5 min. To the resultant mixture solution of N-Fluorobenzenesulfonimide (NFSi) in dry THF (9.58 g of NFSi reagent dissolved in 30 mL of dry THF, 30.4 mmol) was added in the increments of 6 mL. Addition of each increment of NFSi solution was followed by addition of 6 mL of 1M THF solution of NaHMDS (to the total volume of 24 mL of NaHMDS solution). During additions the formation of creamy brown precipitate was observed. Reaction was stirred for additional hour and the precipitate was filtered off and washed with hexane. The filtrate was concentrated *in vacuo* to yield brown oil which dissolved in dichloromethylene enad washed with 1M aqueous solution of sodium bicarbonate. Organic layer was dried over magnesium sulphate, filtered, concentrated *in vacuo* and subjected to column chromatography on silica (chloroform/ ethyl acetate, 0-50%). Product was eluted with 15% ethyl acetate, followed by monofluorination product and unreacted substrate. The tetraisopropyl difluorobisphosphonate was obtained as a pale yellow oil (1.32 g, 3.5 mmol, 43%).

The obtained tetraisopropyl difluorobisphosphonate (1.32 g, 3.5 mmol) was dissolved in dichloromethylene (5 mL) and then transferred to the flask fitted with reflux condenser with tube filled with calcium chloride. To resulting solution TMSBr was added (2.3 mL, 17.4 mmol) and mixture was refluxed for 16 h. Afterwards flask was cooled down to the room temperature and 2.5 mL of methanol was added dropwise. Resultant brown solution was evaporated with two portions (10 mL) of methanol and then treated with 25 mL of water. Mixture was extracted with ethyl acetate until aqueous solution become colourless. The trimethylamine was added (0.98 mL, 7 mmol) to the aqueous phase and resulting mixture was evaporated *in vacuo* to yield pale brown glassy solid.

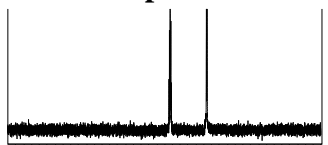
¹⁹F NMR (376 MHz, D₂O): δ -121.42 (2F, t, J = 83.7 Hz); **³¹P NMR** (162 MHz, D₂O) δ 3.44 (2P, t, J = 83.8 Hz). **HRMS** (ESI) calc. for CH₃F₂O₆P₂⁻ requires 210.9378, found 210.9371.

¹H NMR spectrum of m⁷GppCCL₂ppG



PPM: 8.1 7.6 6.5 5.4 4.1 3.1 2.3 1.6 0.8
file: /usr/local/chem/1h/1h_nmr/02390082700001.expt: "s2pul"
Date: 2009-09-24 10:20:09 ppm: 2.3119947 cm/pt: 38563
Number of scans: 64

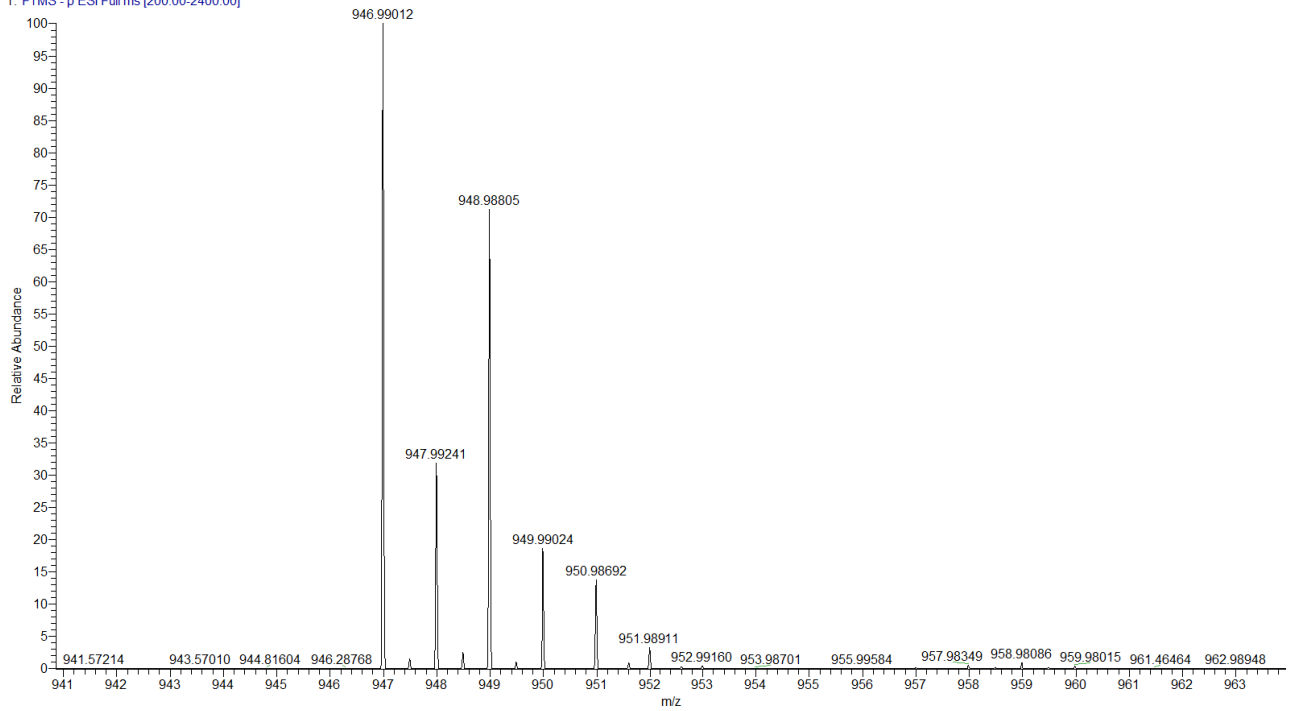
³¹P NMR spectrum of m⁷GppCCL₂ppG



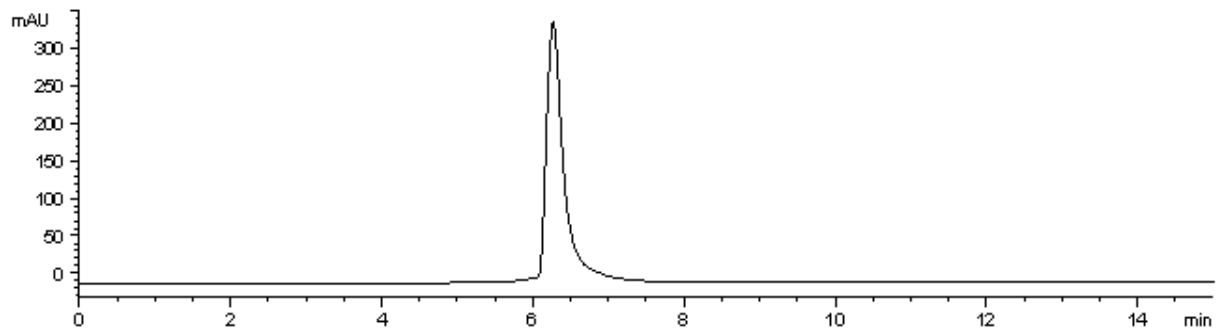
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Number of scans: 128

HRMS spectrum of m⁷GppCCl₂ppG

50526_MBH_07 #5-238 RT: 0.02-1.06 AV: 234 NL: 5.48E5
T: FTMS - p ESI Full ms [200.00-2400.00]

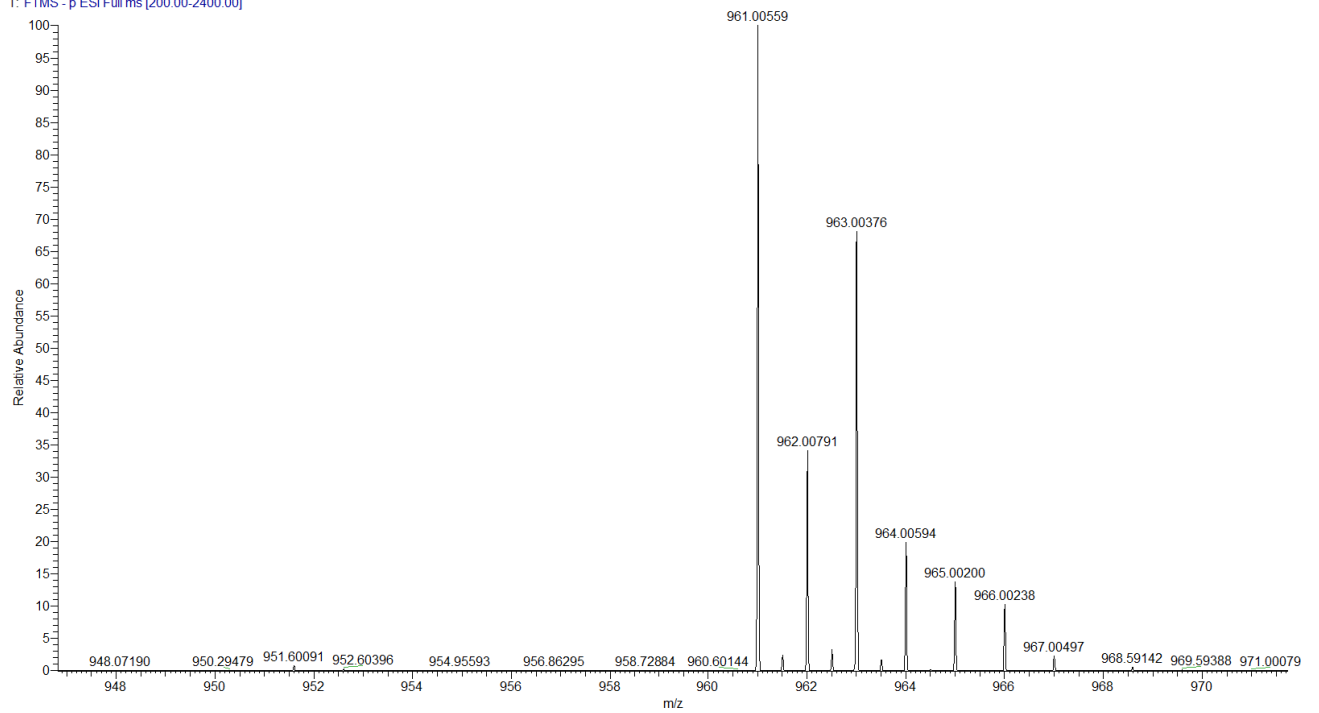


HPLC profile of m⁷GppCCl₂ppG

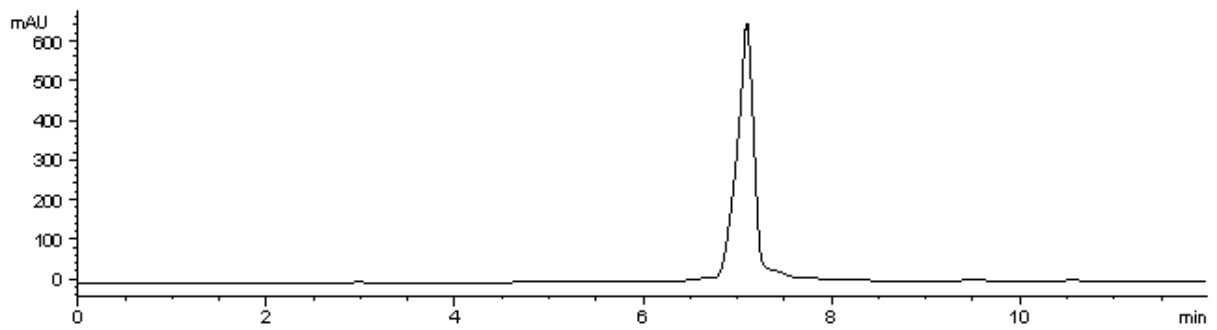


HRMS spectrum of $m_2^{7,2'-O}$ GppCCl₂ppG

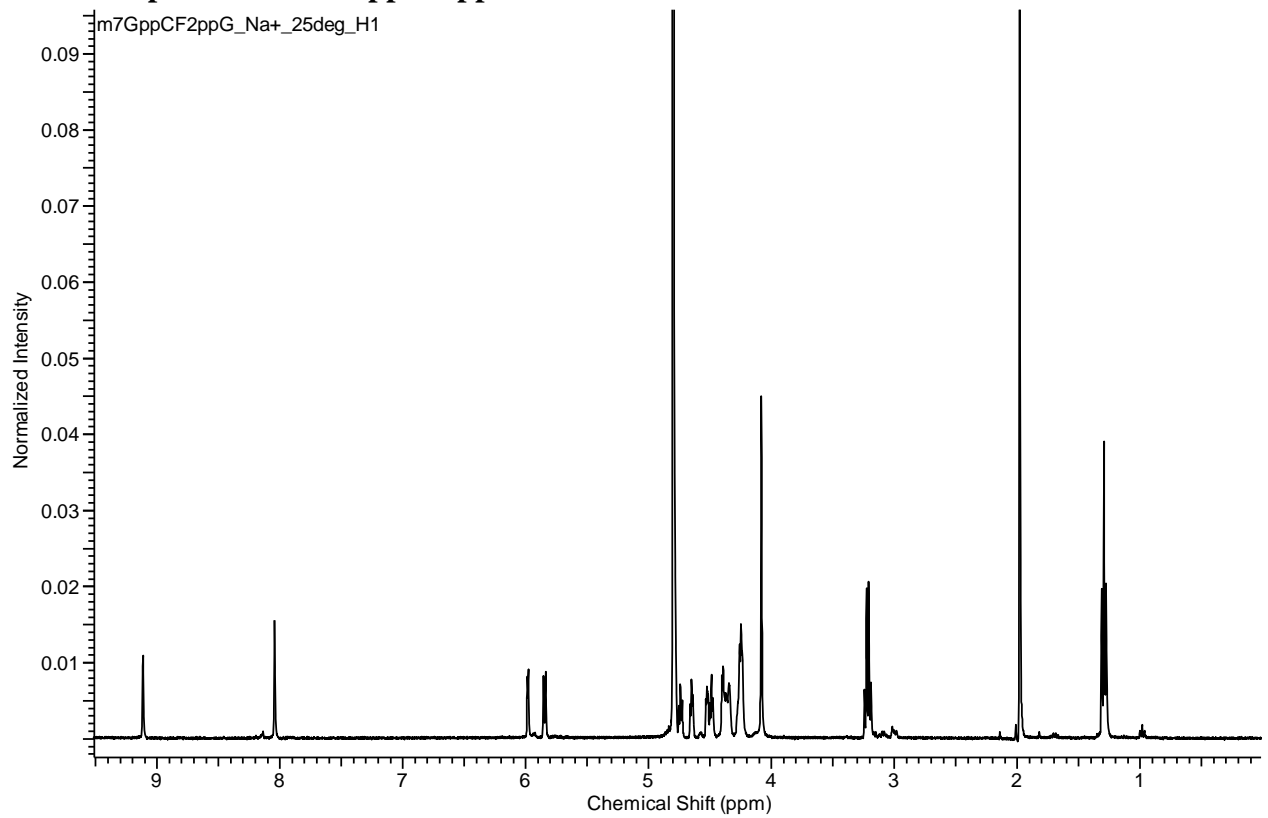
50526_MBH_03 #6-193 RT: 0.03-0.86 AV: 188 NL: 1.38E6
T: FTMS - p ESI Full ms [200.00-2400.00]



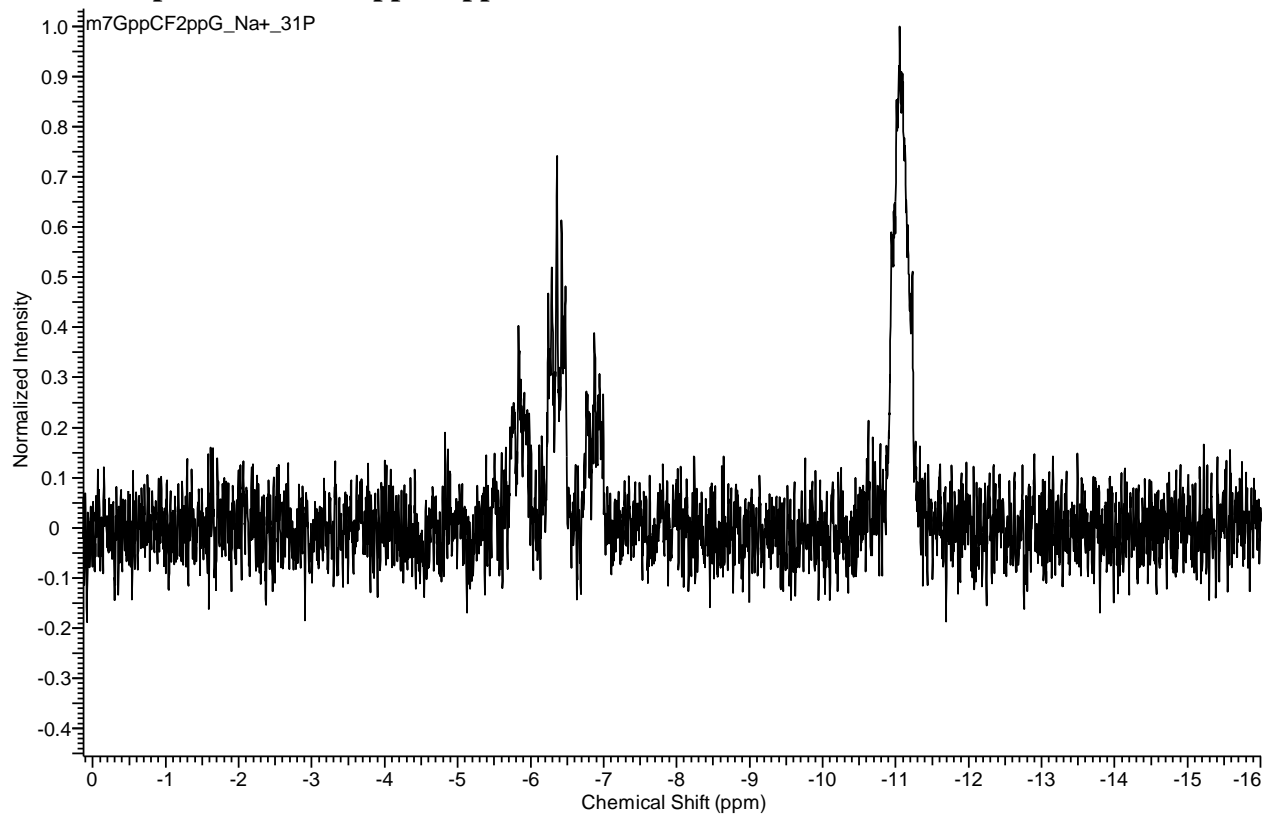
HPLC profile of $m_2^{7,2'-O}$ GppCCl₂ppG



^1H NMR spectrum of $\text{m}^7\text{GppCF}_2\text{ppG}$

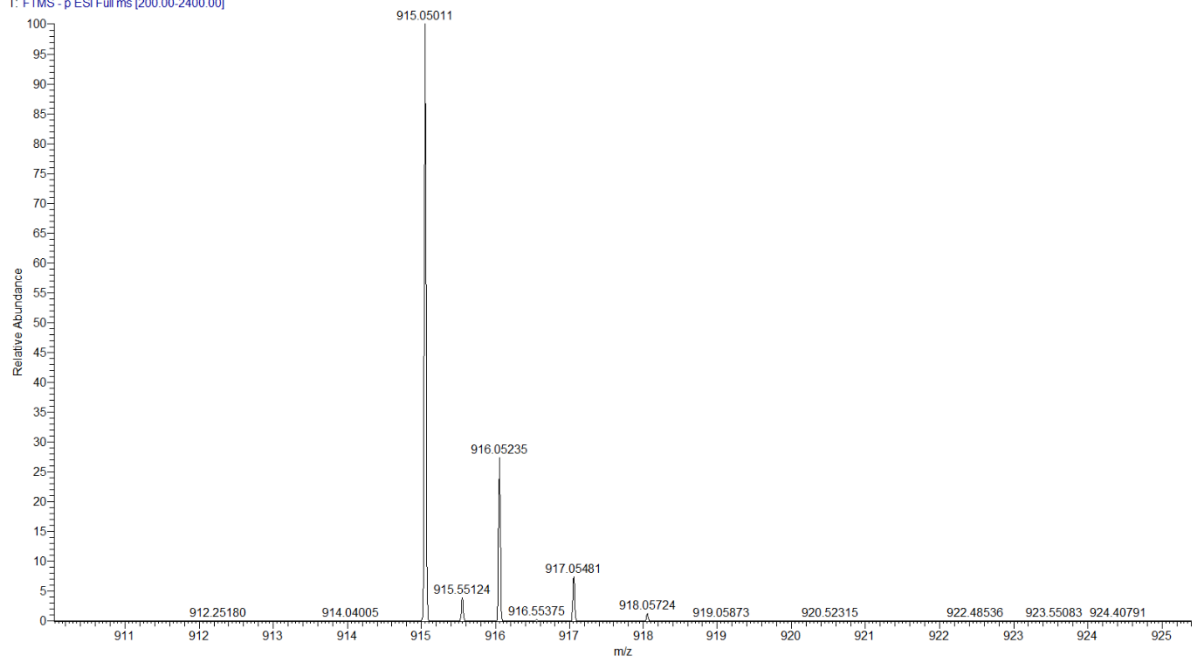


^{31}P NMR spectrum of $\text{m}^7\text{GppCF}_2\text{ppG}$

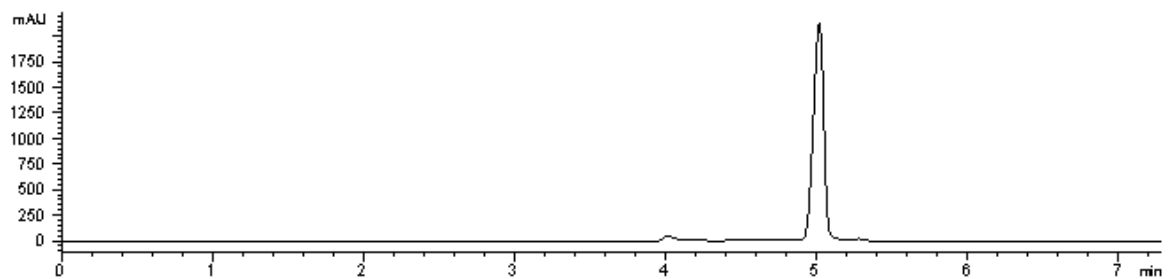


HRMS spectrum of m⁷GppCF₂ppG

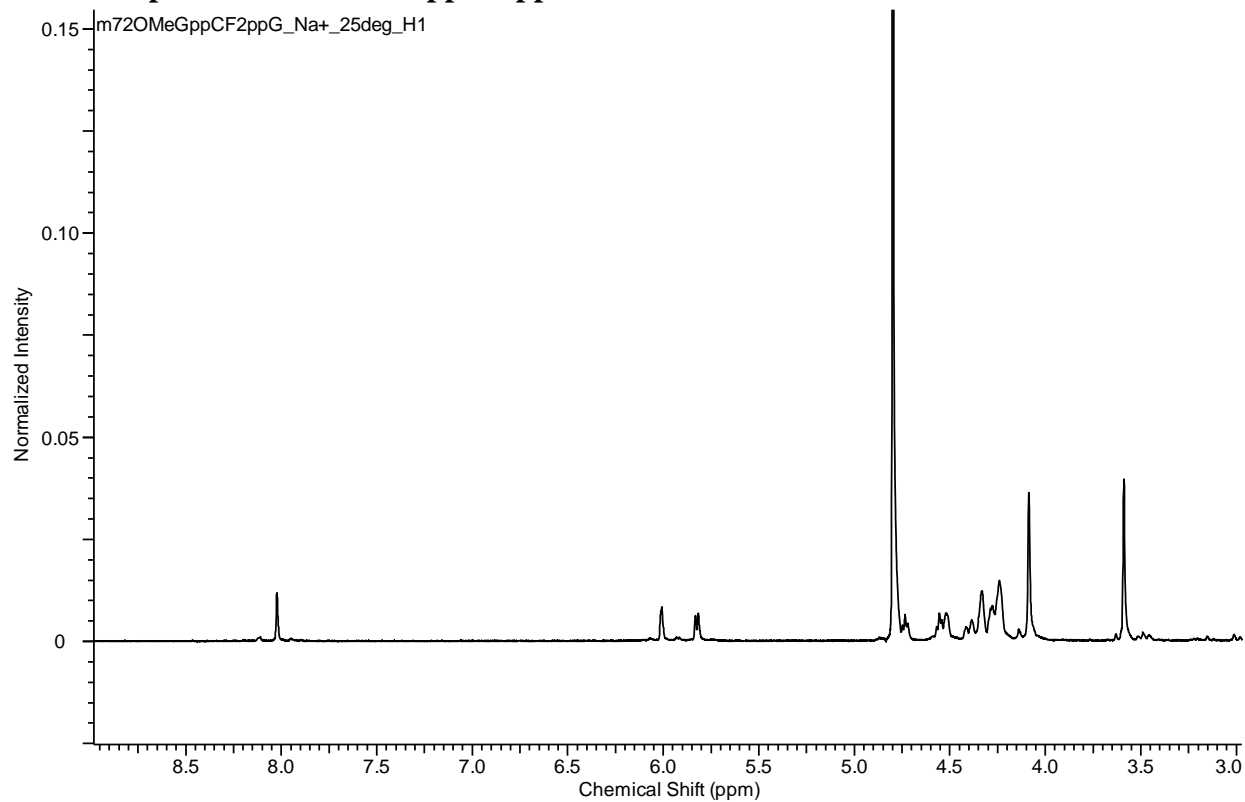
50526_MBH_05 #47-127 RT: 0.21-0.57 AV: 81 NL: 1.64E6
T: FTMS - p ESI Full ms [200.00-2400.00]



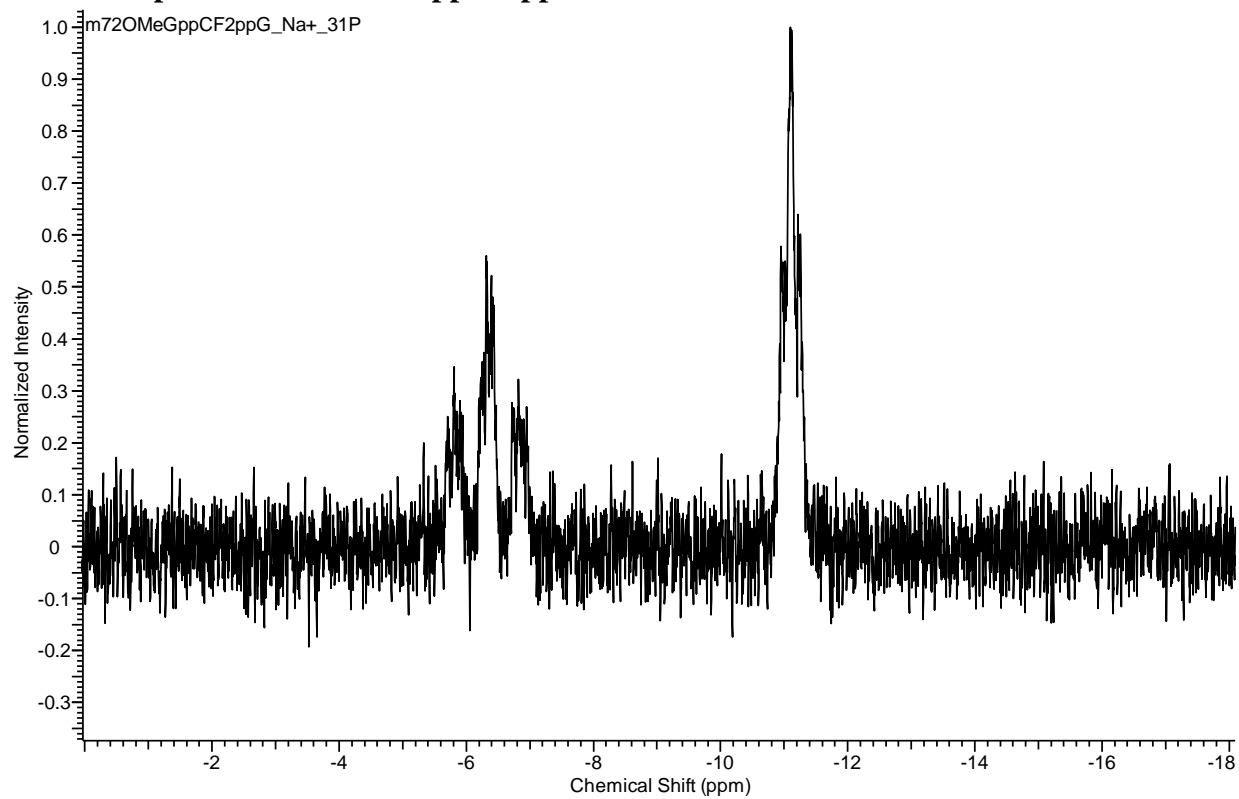
HPLC profile of m⁷GppCF₂ppG



^1H NMR spectrum of $m_2^{7,2'}\text{-O}^-\text{GppCF}_2\text{ppG}$

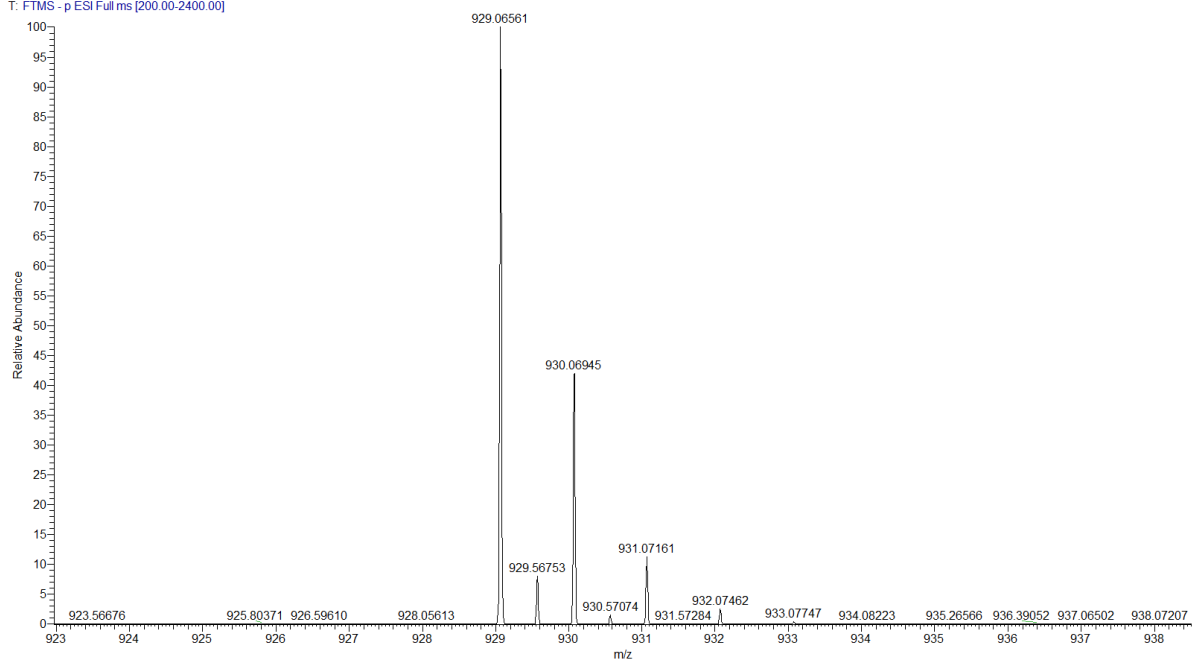


^{31}P NMR spectrum of $m_2^{7,2'}\text{-O}^-\text{GppCF}_2\text{ppG}$

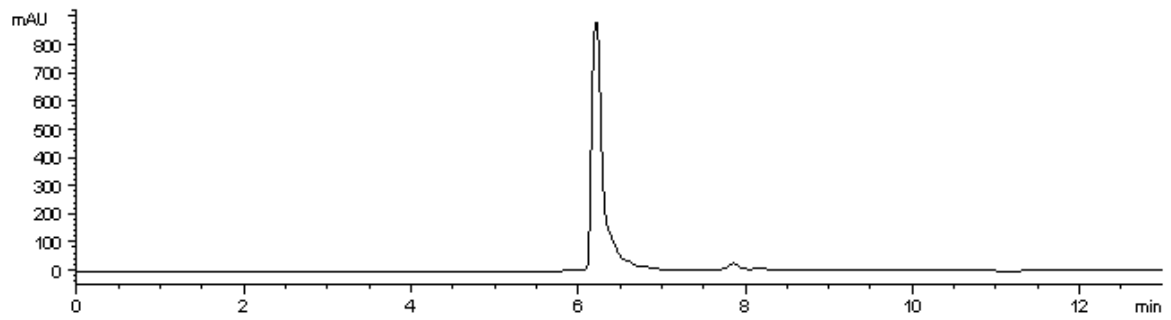


HRMS spectrum of $m_2^{7,2'-O}GppCF_2ppG$

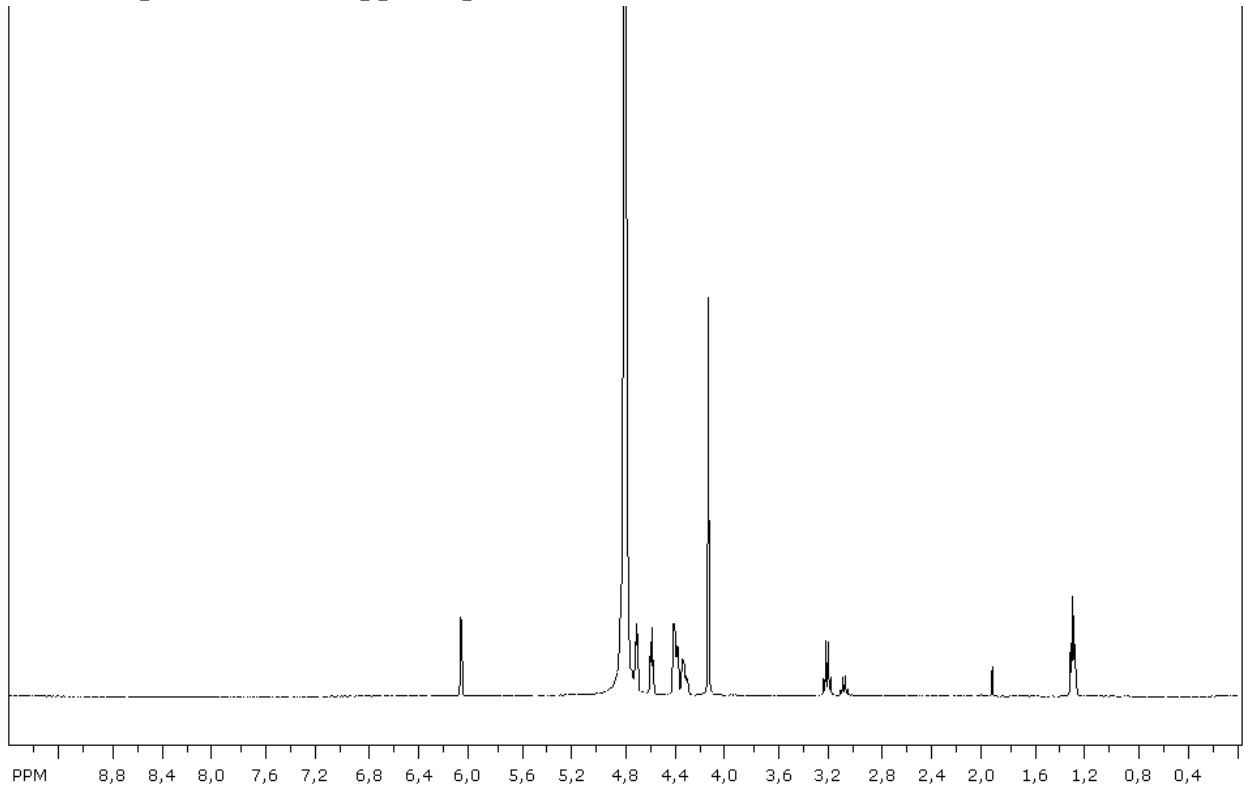
50526_MBH_08_#47-179 RT: 0.21-0.80 AV: 133 NL: 3.83E6
T: FTMS - p ESI Full ms [200.00-2400.00]



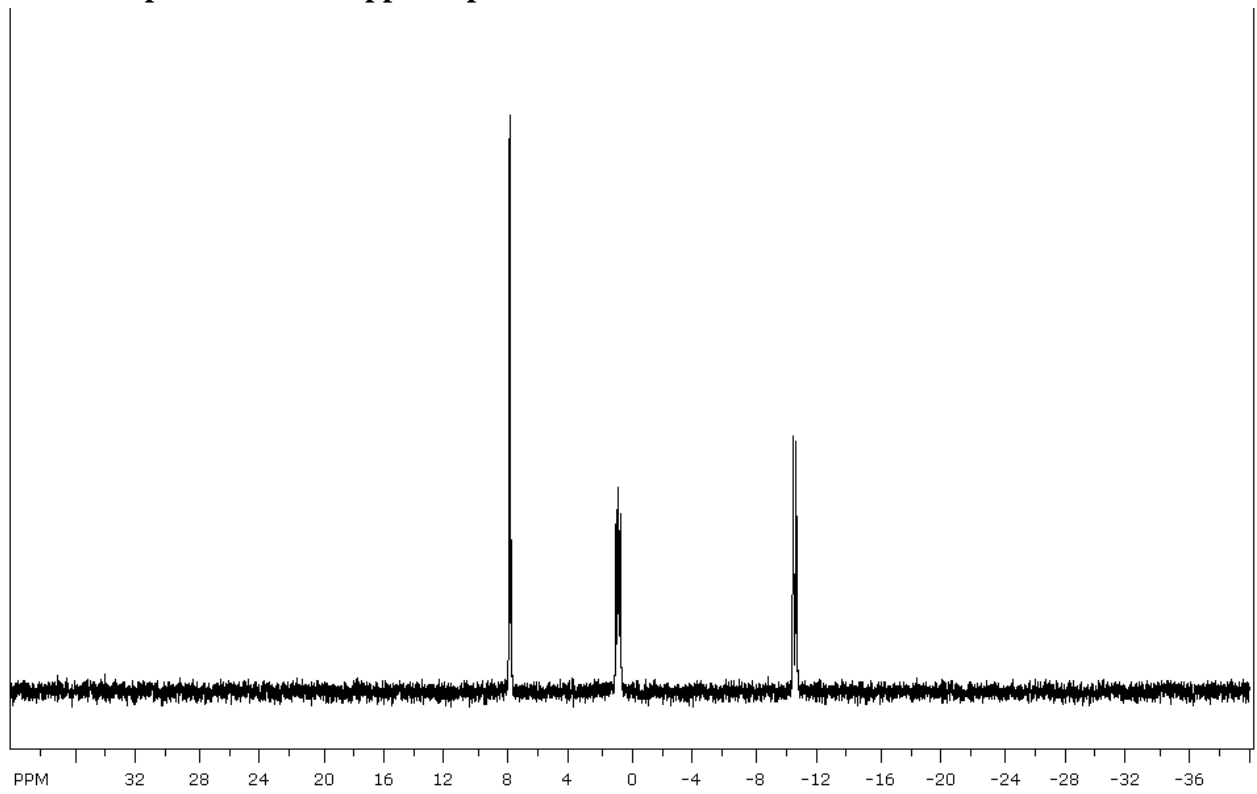
HPLC profile of $m_2^{7,2'-O}GppCF_2ppG$



^1H NMR spectrum of $m^7\text{GppCCl}_2\text{p}$

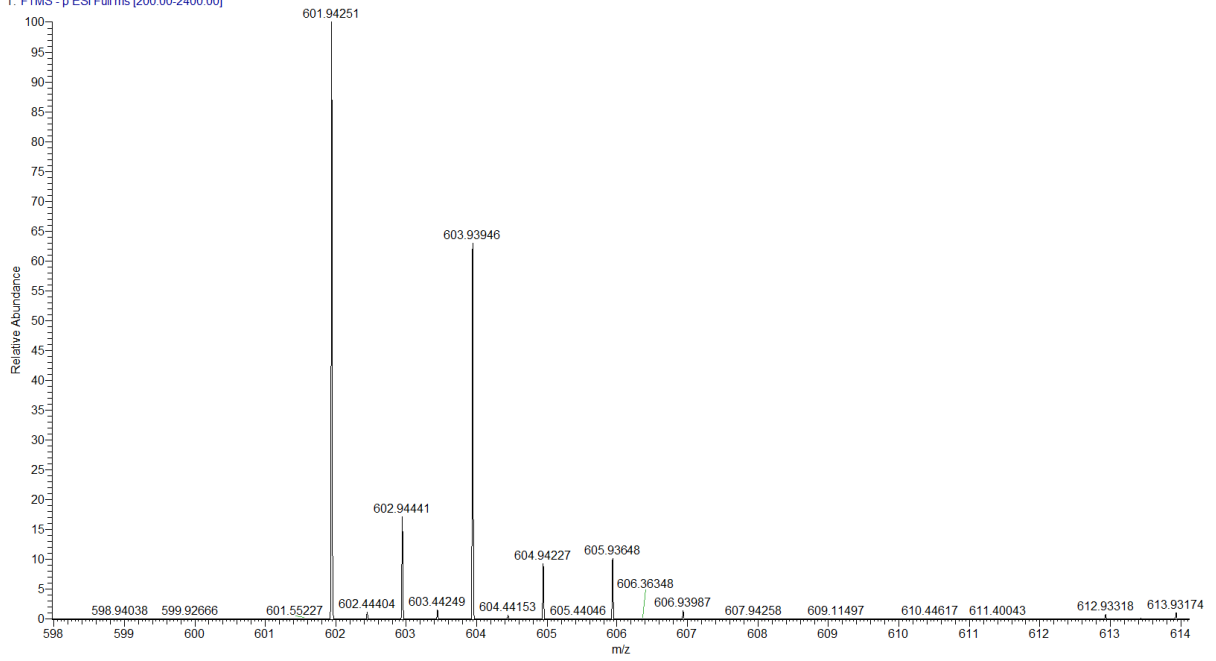


^{31}P NMR spectrum of $m^7\text{GppCCl}_2\text{p}$

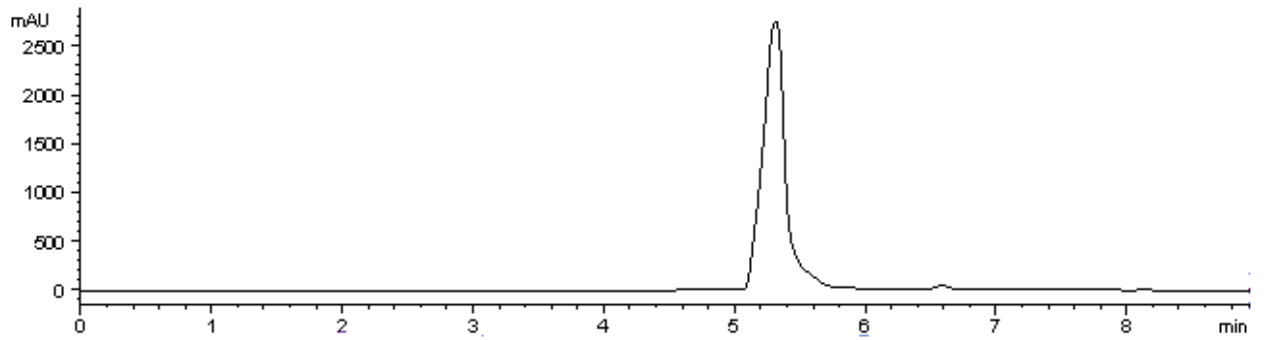


HRMS spectrum of m⁷GppCCl₂p

50526_MBH_02_#43-472 RT: 0.19-2.10 AV: 430 NL: 6.65E5
T: FTMS - p ESI Full ms [200.00-2400.00]

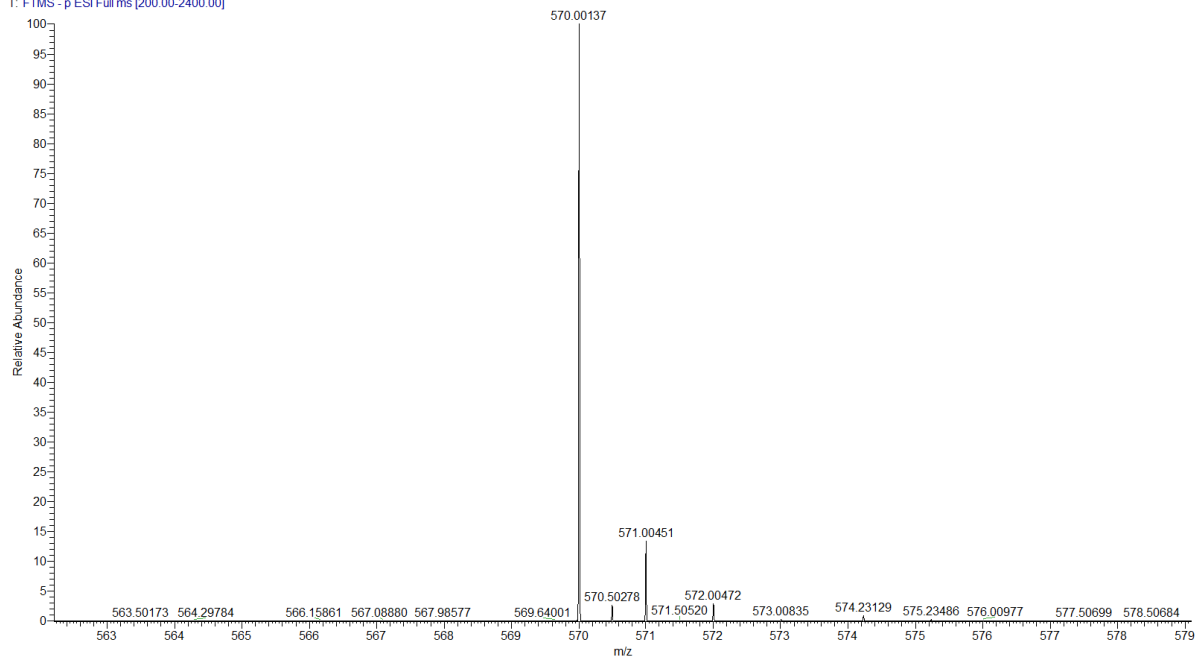


HPLC profile of m⁷GppCCl₂p

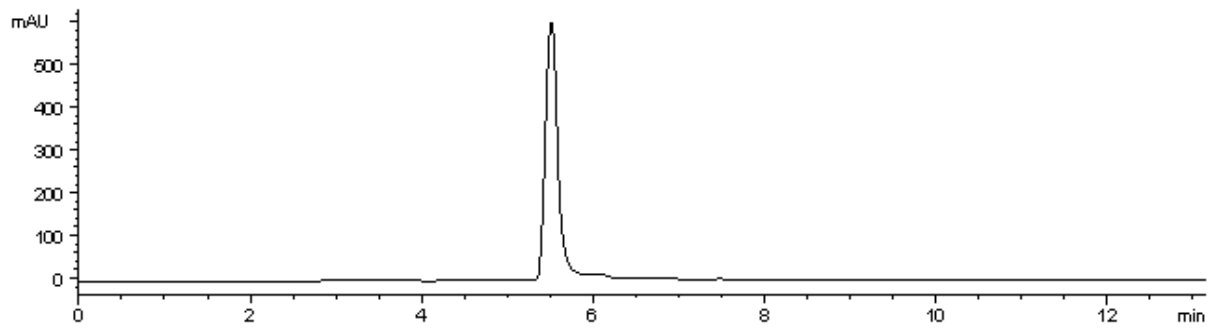


HRMS spectrum of m⁷GppCF₂p

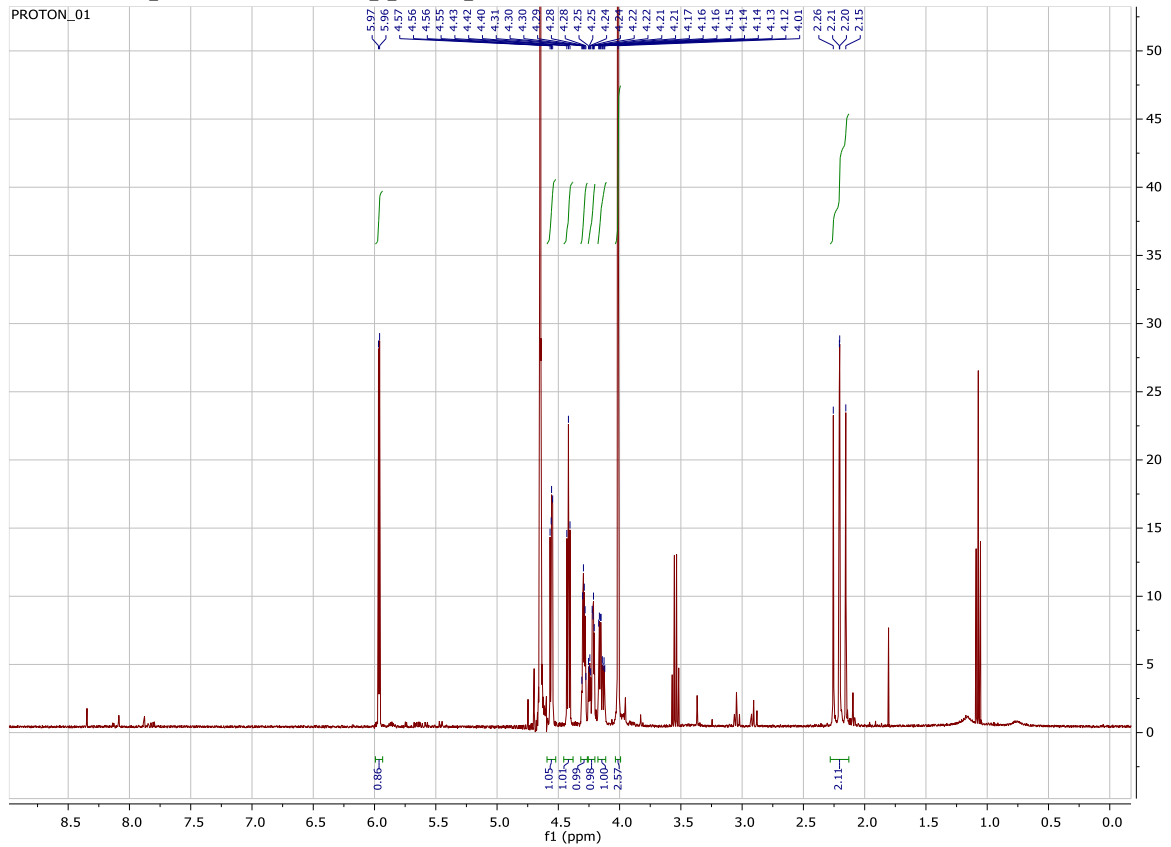
50526_MBH_06_#47-182_RT: 0.21-0.81_AV: 136_NL: 2.58E6
T: FTMS - p ESI Full ms [200.00-2400.00]



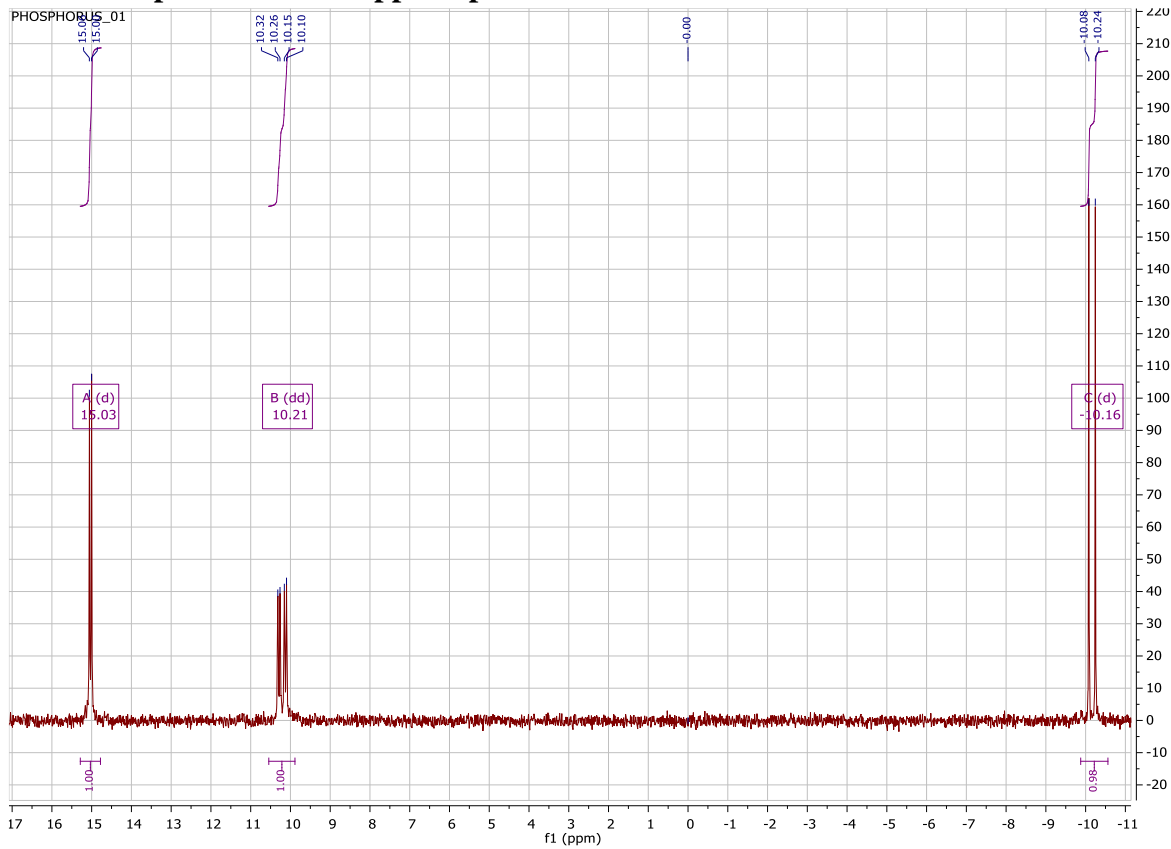
HPLC profile of m⁷GppCF₂p



^1H NMR spectrum of $m^7\text{GppCH}_2\text{p}$

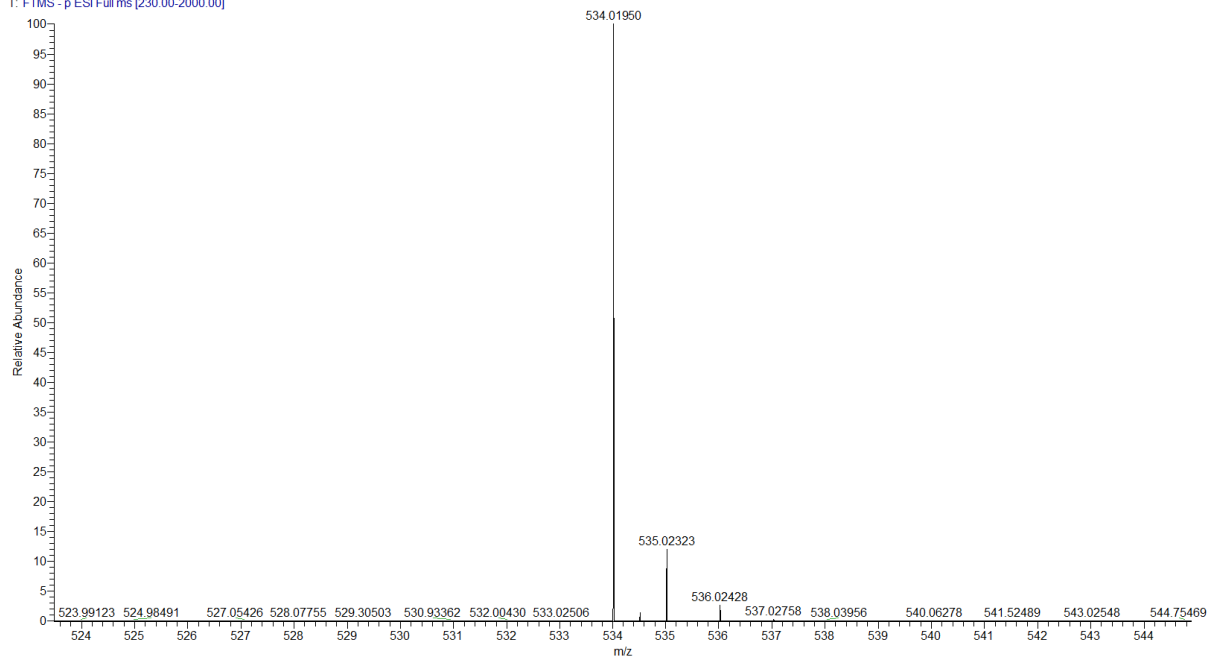


^{31}P NMR spectrum of $m^7\text{GppCH}_2\text{p}$

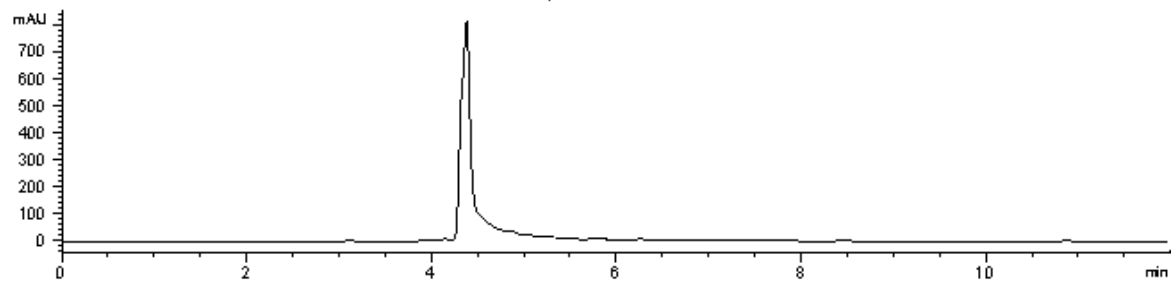


HRMS spectrum of m⁷GppCH₂p

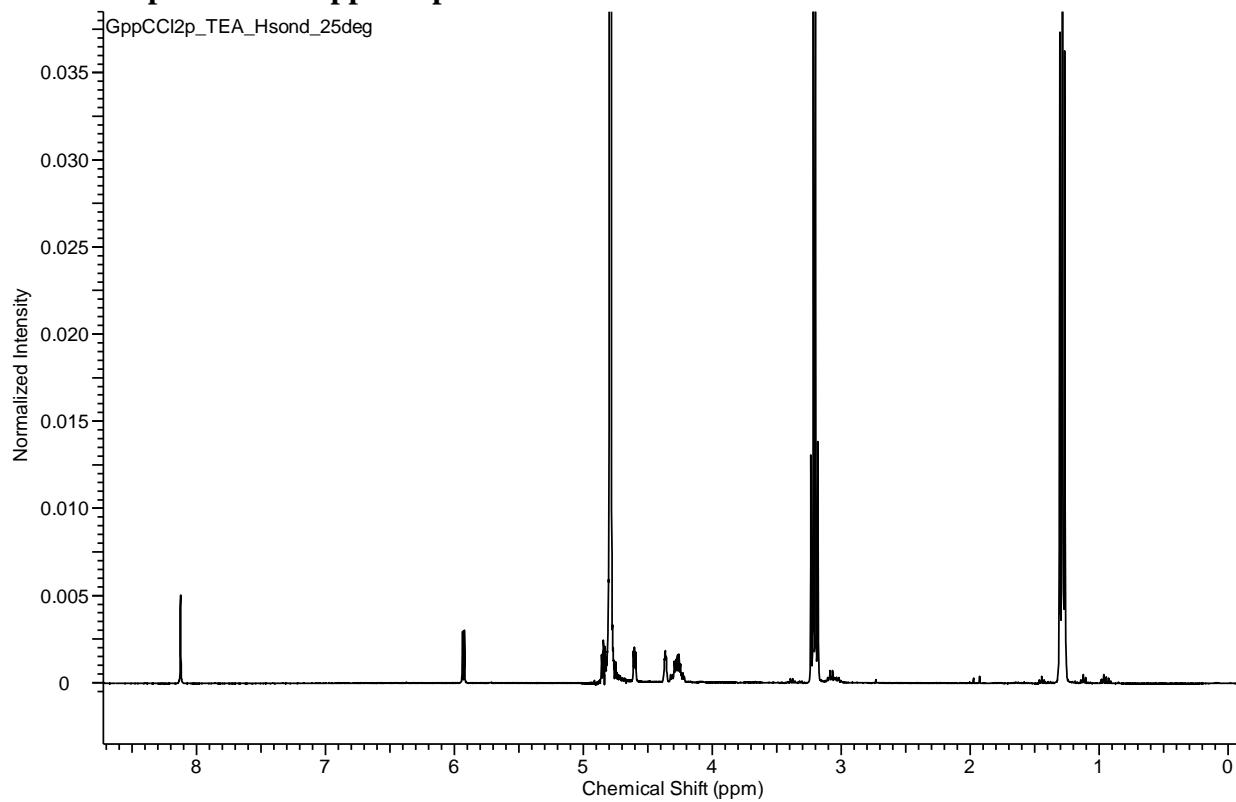
51105_MBAR2 #53-163 RT: 0.52-1.81 AV: 131 NL: 8.94E6
T: FTMS - p ESI Full ms [230.00-2000.00]



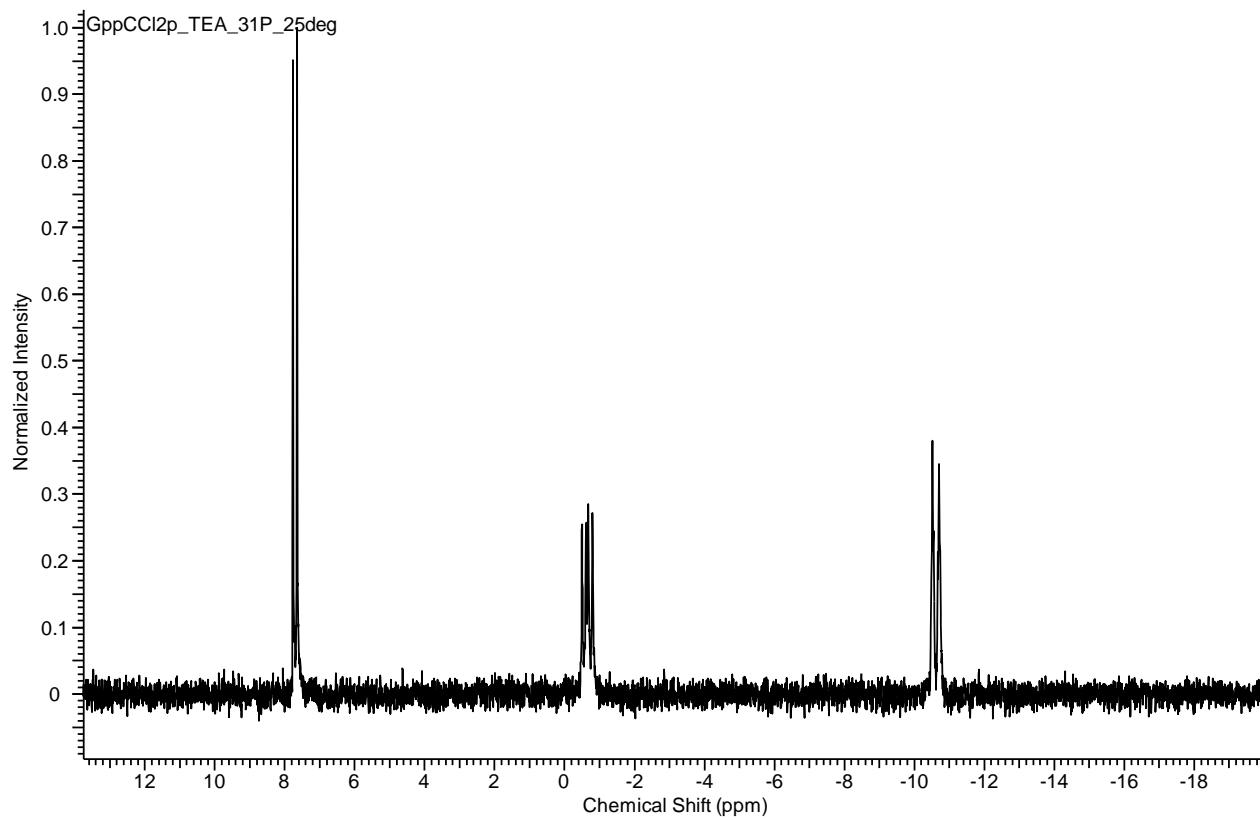
HPLC profile of m⁷GppCH₂p



¹H NMR spectrum of GppCCl₂p

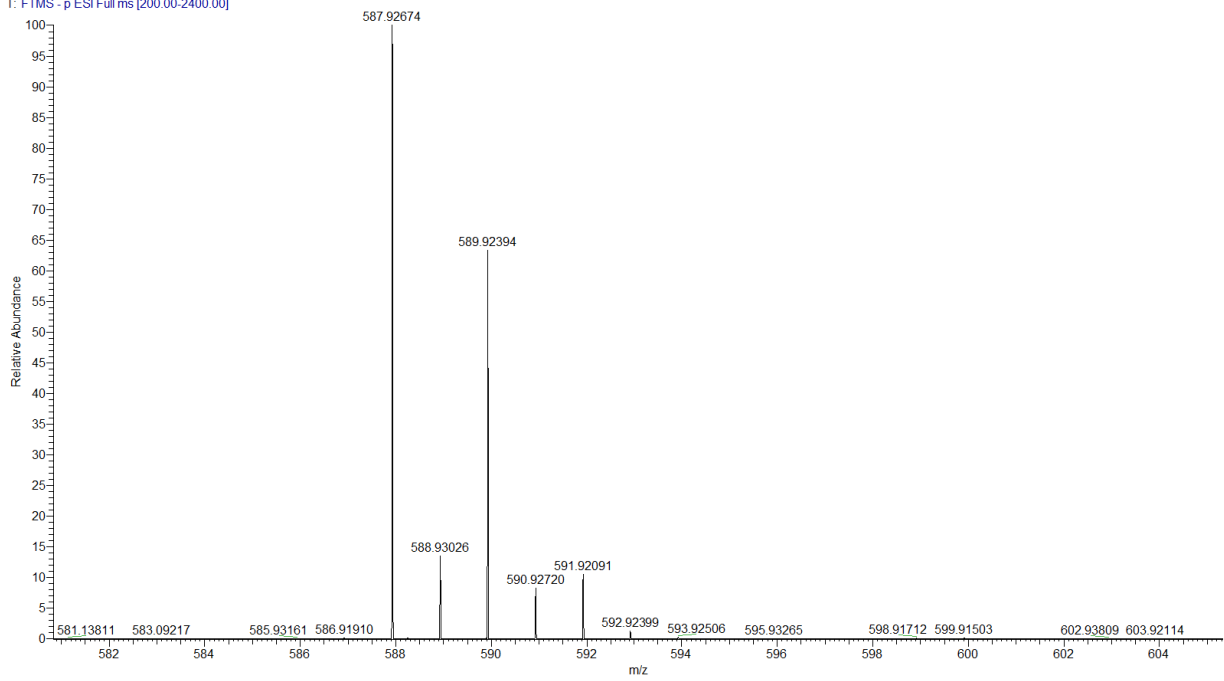


³¹P NMR spectrum of GppCCl₂p

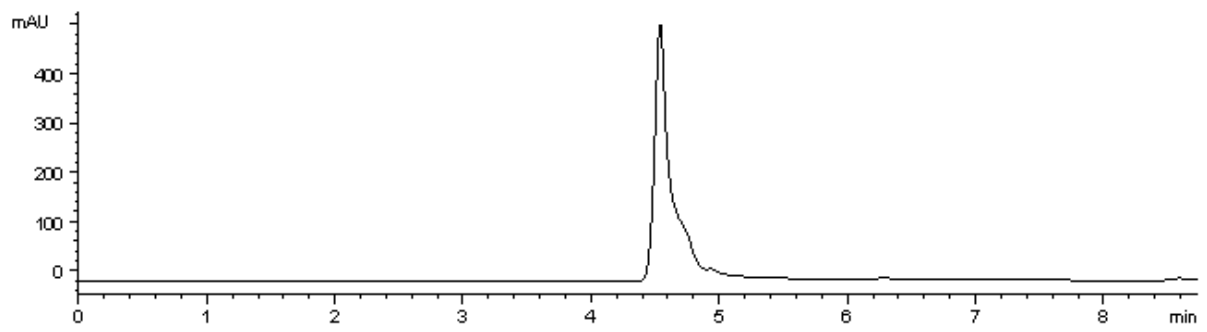


HRMS spectrum of GppCCl₂p

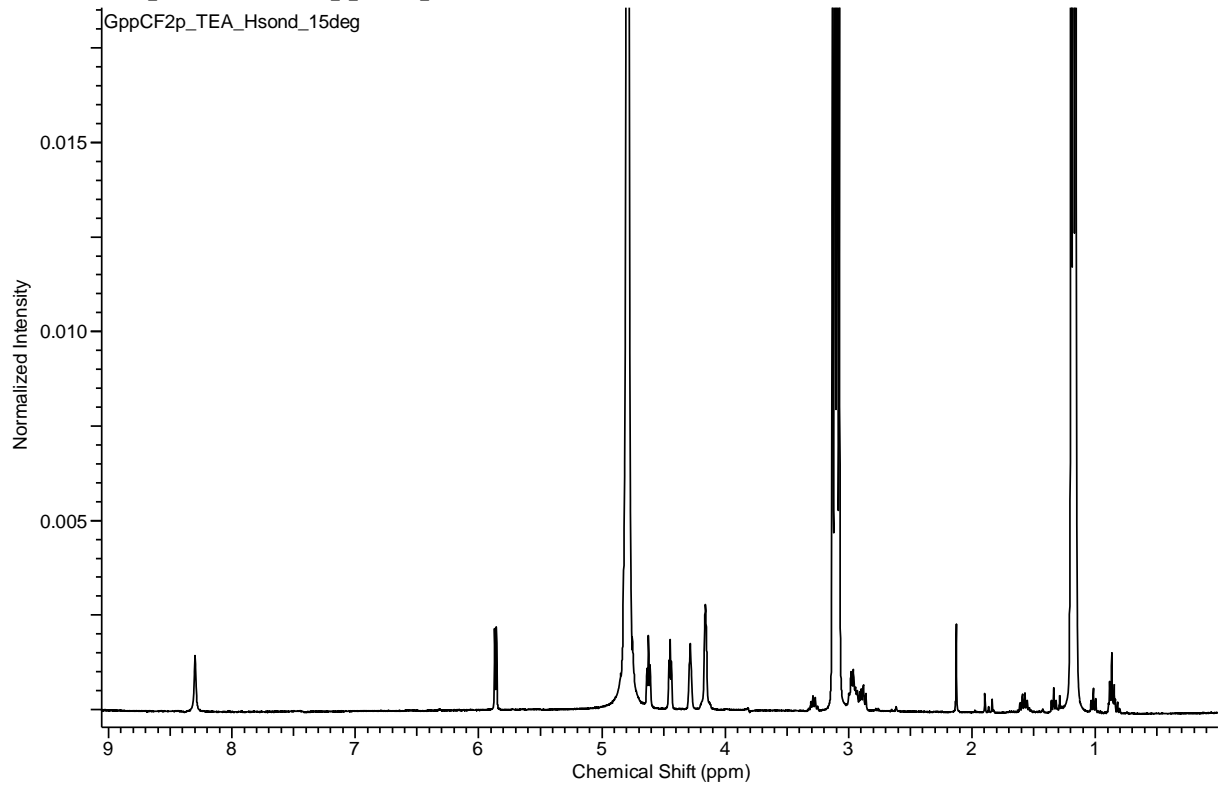
50526_MBH_04 #61-293 RT: 0.27-1.31 AV: 233 NL: 6.12E6
T: FTMS - p ESI Full ms [200.00-2400.00]



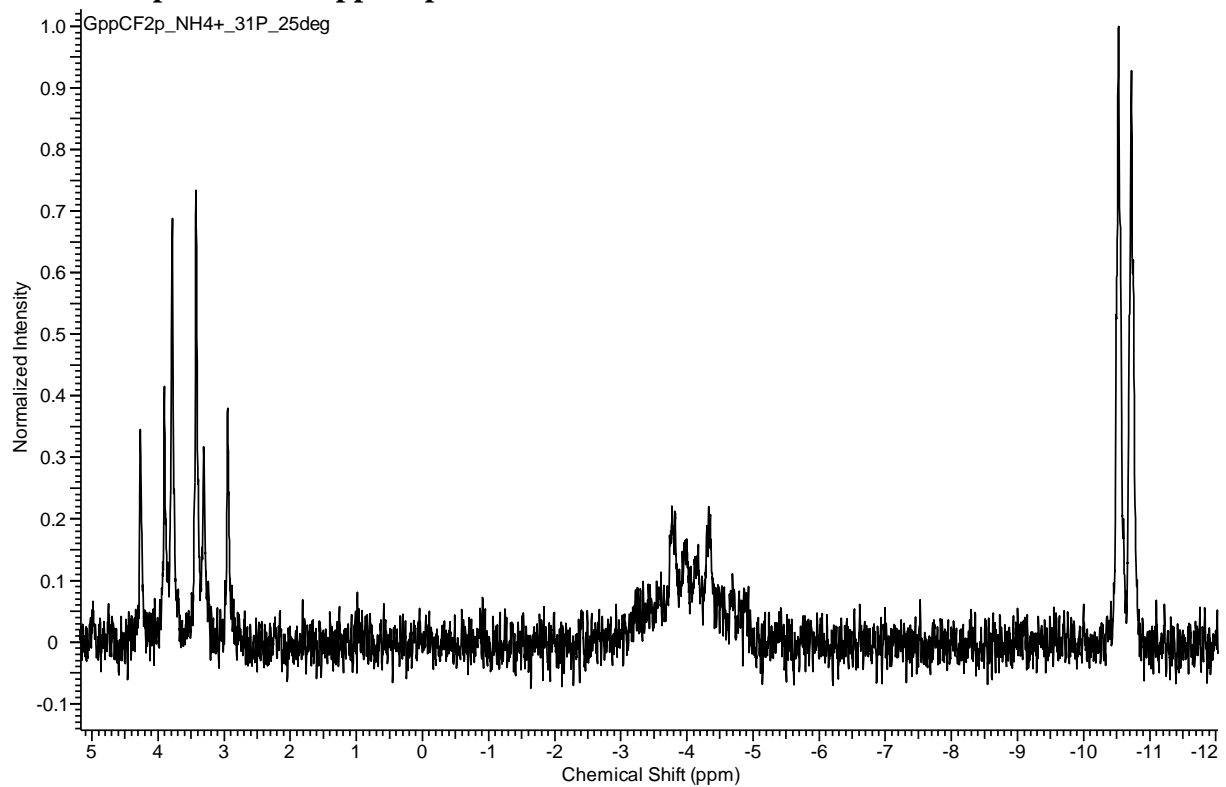
HPLC profile of GppCCl₂p



¹H NMR spectrum of GppCF₂p

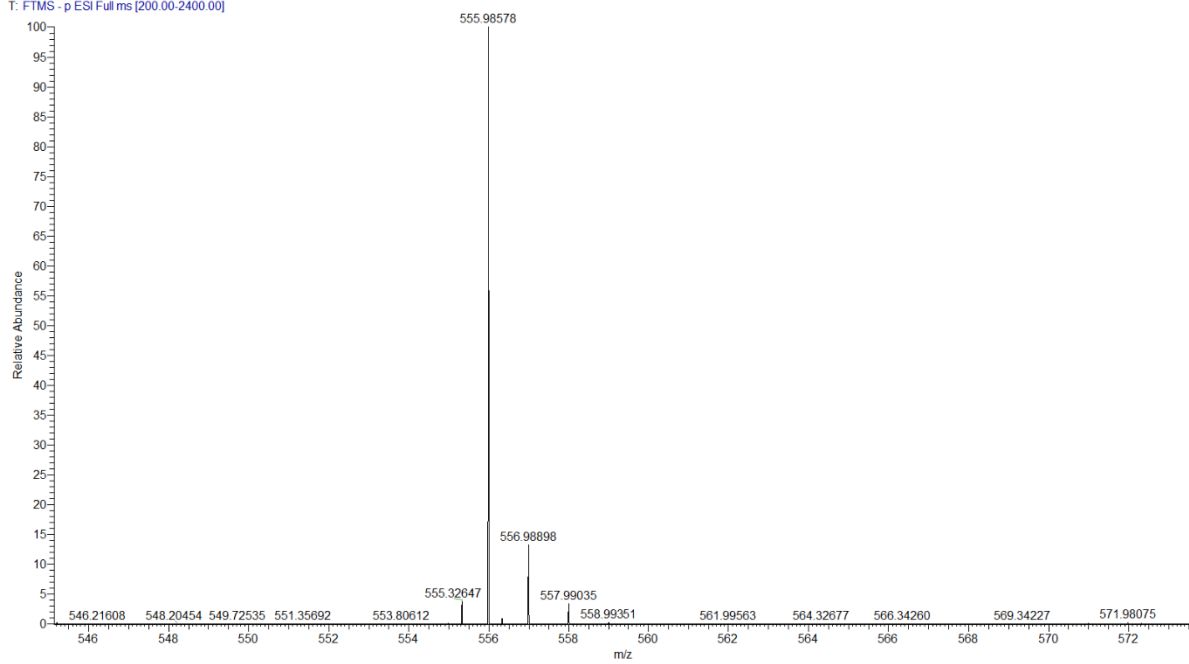


³¹P NMR spectrum of GppCF₂p

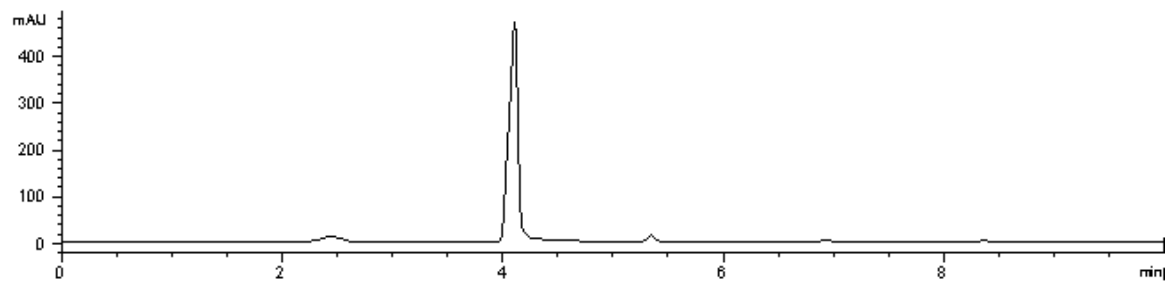


HRMS spectrum of GppCF₂p

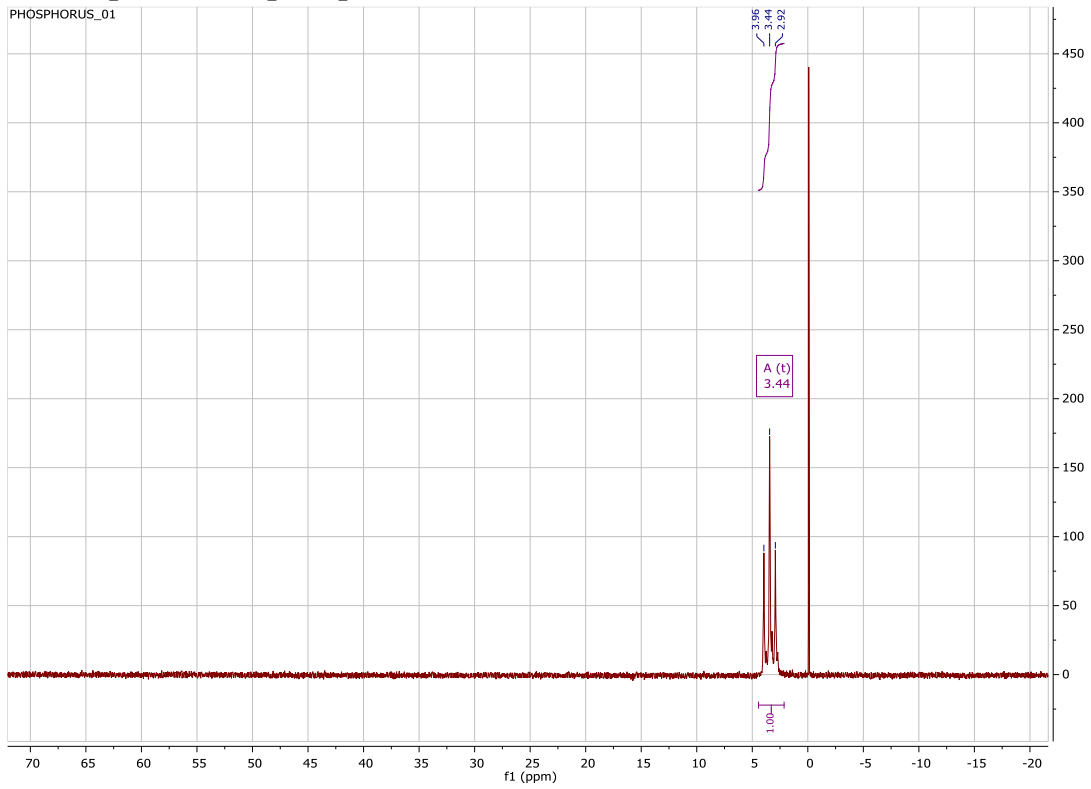
50528_MBH_01#710-1133 RT: 3.16-5.05 AV: 424 NL: 8.93E5
T: FTMS - p ESI Full ms [200.00-2400.00]



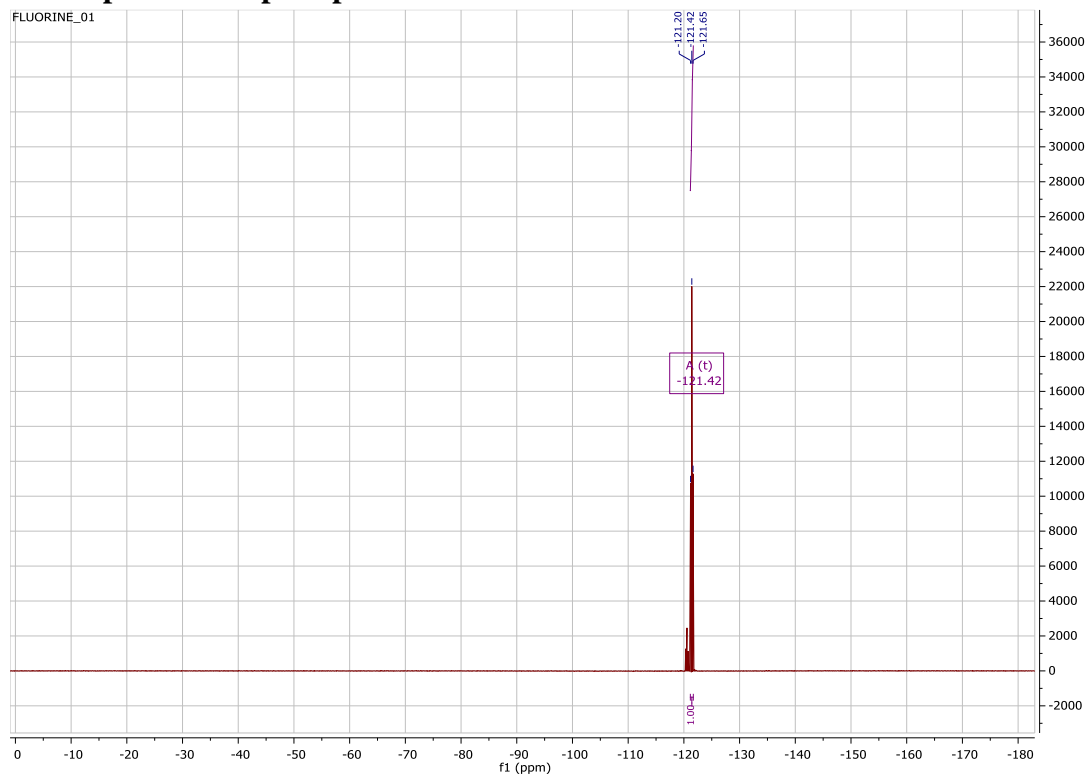
HPLC profile of GppCF₂p



³¹P NMR spectrum of pCF₂p

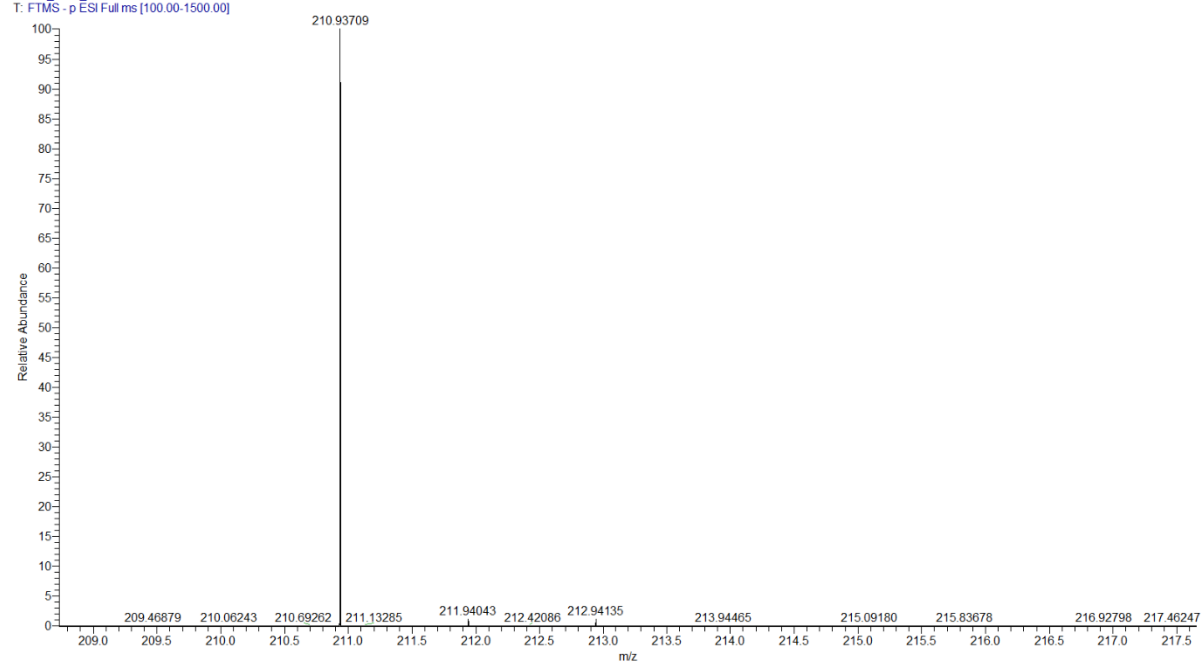


¹⁹F NMR spectrum of pCF₂p



HRMS spectrum of pCF₂p

50526_MBH_09 #33-147 RT: 0.15-0.66 AV: 115 NL: 4.18E8
T: FTMS - p ESI Full ms [100.00-1500.00]



References

- [1] Blackburn, G. M., Kent, D. E., and Kolkman, F. (1981) Three new β , γ -methylene analogues of adenosine triphosphate, *J. Chem. Soc. Chem. Comm.*, 1188-1190.
- [2] Rydzik, A. M., Lukaszewicz, M., Zuberek, J., Kowalska, J., Darzynkiewicz, Z. M., Darzynkiewicz, E., and Jemielity, J. (2009) Synthetic dinucleotide mRNA cap analogs with tetraphosphate 5',5' bridge containing methylenebis(phosphonate) modification, *Org. Biomol. Chem.* 7, 4763-4776.
- [3] Rydzik, A. M., Lukaszewicz, M., Zuberek, J., Kowalska, J., Darzynkiewicz, Z. M., Darzynkiewicz, E., and Jemielity, J. (2009) Synthetic dinucleotide mRNA cap analogs with tetraphosphate 5',5' bridge containing methylenebis(phosphonate) modification, *Org. Biomol. Chem.* 7, 4763-4776.
- [4] Jemielity, J., Fowler, T., Zuberek, J., Stepinski, J., Lewdorowicz, M., Niedzwiecka, A., Stolarski, R., Darzynkiewicz, E., and Rhoads, R. E. (2003) Novel "anti-reverse" cap analogs with superior translational properties, *RNA* 9, 1108-1122.
- [5] Jemielity, J., Stepinski, J., Jaremko, M., Haber, D., Stolarski, R., Rhoads, R. E., and Darzynkiewicz, E. (2003) Synthesis of Novel mRNA 5' Cap-Analogues: Dinucleoside P1, P3-Tri-, P1, P4-Tetra-, and P1, P5-Pentaphosphates, *Nucleosides, Nucleotides and Nucleic Acids* 22, 691-694.

